Leu Pro Glu Leu Lys Gly Lys Glu Lys Leu Gly Glu Thr Ser Leu Phe 170 165

Val Leu Val Lys Ser Ala Ala 180

- (2) INFORMATION FOR SEQ ID NO:1170:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 870 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..870
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597162
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1170: atggcgactc tcttcactgc gacggttcct tctcaccacc gtttcgtctc tccatcccaa 60 catccgaaac agagtettet ategeagteg etaagegtea cattcaegga gaatecaeaa 120 ccaacggcgg tggtcacatt gcaggagcag caattgacgg attggattac ttctccggtg 180 acgcgacggt toggaatogg ogcogggttt acctgggctg ggtttttagc tttcggtgtt 240 gtctccgagc agatgaagaa gtcacgactc gatgtgtttc aggaagagga taacacaaga 300 ggtttagaaa agcaagaaga gatcatctta ccaaacggca taaggtacta tgatctacaa 360 gttggaagtg gagctactcc aagctcagga tacttggtag tgtttgatgt aaagggacaa 420 gtacatggca ctgaacaagt gticgtggac acattiggag gcaaaggcaa gtcactagcg 480 atggtaatgg actcaagacc gtatagcaag ggactatgcc aaggtataga gcatgttctc 540 aggtcaatga aggctggagg taaacgtaga gtgataattc caccgtcctt aggatttgga 600 gacagaaatg togaatttgg acagggtttg gagatteete catetgeaac acttgactat 660 atcatcgagg ttgatacagt ttattgtttc caaactattg tggtttacca gttcatattc 720 ttcgtaatca cggctctttt caaatttgat caagtcactg actttgctgg tagcacaaac 780 ttegttatae ttgetgtgtt aacaettgtt etcaaageet ettggeattt tegacaggat 840 cttgcaatgg ggggaagatc gtcgctttga
- (2) INFORMATION FOR SEQ ID NO:1171:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 289 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..289
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597163
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1171:
- Met Ala Thr Leu Phe Thr Ala Thr Val Pro Ser His His Arg Phe Val 10 Ser Pro Ser Gln His Pro Lys Gln Ser Leu Leu Ser Gln Ser Leu Ser
 - 25
- Val Thr Phe Thr Glu Asn Pro Gln Pro Thr Ala Val Val Thr Leu Gln 40 Glu Gln Gln Leu Thr Asp Trp Ile Thr Ser Pro Val Thr Arg Arg Phe
- 60 55 Gly Ile Gly Ala Gly Phe Thr Trp Ala Gly Phe Leu Ala Phe Gly Val
- 75 70 Val Ser Glu Gln Met Lys Lys Ser Arg Leu Asp Val Phe Gln Glu Glu 90
- 85 Asp Asn Thr Arg Gly Leu Glu Lys Gln Glu Glu Ile Ile Leu Pro Asn 110 105 100
- Gly Ile Arg Tyr Tyr Asp Leu Gln Val Gly Ser Gly Ala Thr Pro Ser 125 120
- Ser Gly Tyr Leu Val Val Phe Asp Val Lys Gly Gln Val His Gly Thr 140 135 130

Len

Glu Gln Val Phe Val Asp Thr Phe Gly Gly Lys Gly Lys Ser Leu Ala 150 155 Met Val Met Asp Ser Arg Pro Tyr Ser Lys Gly Leu Cys Gln Gly Ile 165 170 Glu His Val Leu Arg Ser Met Lys Ala Gly Gly Lys Arg Arg Val Ile 185 180 Ile Pro Pro Ser Leu Gly Phe Gly Asp Arg Asn Val Glu Phe Gly Gln 205 195 200 Gly Leu Glu Ile Pro Pro Ser Ala Thr Leu Asp Tyr Ile Ile Glu Val 215 220 Asp Thr Val Tyr Cys Phe Gln Thr Ile Val Val Tyr Gln Phe Ile Phe 235 230 Phe Val Ile Thr Ala Leu Phe Lys Phe Asp Gln Val Thr Asp Phe Ala 250 2.4.5 Gly Ser Thr Asn Phe Val Ile Leu Ala Val Leu Thr Leu Val Leu Lys 265 270 Ala Ser Trp His Phe Arg Gln Asp Leu Ala Met Gly Gly Arg Ser Ser 285 280

- (2) INFORMATION FOR SEQ ID NO:1172:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 205 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..205
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597165
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1172: Met Lys Lys Ser Arg Leu Asp Val Phe Gln Glu Glu Asp Asn Thr Arg
- 1 5 10 15
 Gly Leu Glu Lys Gln Glu Glu Ile Ile Leu Pro Asn Gly Ile Arg Tyr
 20 25 30
- Tyr Asp Leu Gln Val Gly Ser Gly Ala Thr Pro Ser Ser Gly Tyr Leu 35 40 45
- Val Val Phe Asp Val Lys Gly Gln Val His Gly Thr Glu Gln Val Phe
 50 55 60
- Val Asp Thr Phe Gly Gly Lys Gly Lys Ser Leu Ala Met Val Met Asp
 65 70 75 80
- Ser Arg Pro Tyr Ser Lys Gly Leu Cys Gln Gly Ile Glu His Val Leu
 85
 90
 95
- Arg Ser Met Lys Ala Gly Gly Lys Arg Arg Val Ile Ile Pro Pro Ser 100 105 110 Leu Gly Phe Gly Asp Arg Asn Val Glu Phe Gly Gln Gly Leu Glu Ile
- Pro Pro Ser Ala Thr Leu Asp Tyr Ile Ile Glu Val Asp Thr Val Tyr
- 130 135 140

 Cys Phe Gln Thr Ile Val Val Tyr Gln Phe Ile Phe Phe Val Ile Thr

 145 150 155 160
- Ala Leu Phe Lys Phe Asp Gln Val Thr Asp Phe Ala Gly Ser Thr Asn 165
- Phe Val Ile Leu Ala Val Leu Thr Leu Val Leu Lys Ala Ser Trp His 180 185 190
- Phe Arg Gln Asp Leu Ala Met Gly Gly Arg Ser Ser Leu 195 200 205
- (2) INFORMATION FOR SEQ ID NO:1173:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 600 base pairs

- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..600
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597170
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1173: atggggaagg gtaaccagtt ggaagaagtc agctacatca acaacaacca gggtggctac 60 aaaggataca acaacttcaa aaccaacaat cccaacctct cctaccgtag caccagcgtt 120 gctaatcctc aggatcaggt gtatcccgtg caacaacaac aaggtcagaa caaacctttt 180 gttctctaca accaaggttt cgttcctaag cagcaatttn tggggaacta ccagccgcca 240 300 ccaccacctg gatttgcaca tcagcaaaac catgatggaa acactgaata ccaaagtccg atacttagag ggacattetg catettette agetecaaca caaacaagee aaactaceca 360 ggcaaagcag ttcagaatct aaaagaatat gctcatgcta ttacactgcg tagtggaaaa 420 gcacttccaa ctagggagga accaaagacg gtcactgagg acagtgaaga tcaagatgga gaggatttaa gtctcgagca accactcgac ntgtcactcg agcanccact cgaccaacca 540 ctcgagcaac cactcgaccg tgctactcaa cctatcttcc cagcagtatc atcaacgtag 600
- (2) INFORMATION FOR SEQ ID NO:1174:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 199 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..199
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597171
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1174: Met Gly Lys Gly Asn Gln Leu Glu Glu Val Ser Tyr Ile Asn Asn Asn

1 5 10 15
Gln Gly Gly Tyr Lys Gly Tyr Asn Asn Phe Lys Thr Asn Asn Pro Asn 20 25 30

Leu Ser Tyr Arg Ser Thr Ser Val Ala Asn Pro Gln Asp Gln Val Tyr 35 40 45

Pro Val Gln Gln Gln Gly Gln Asn Lys Pro Phe Val Leu Tyr Asn 50 55 60

Gln Gly Phe Val Pro Lys Gln Gln Phe Xaa Gly Asn Tyr Gln Pro Pro 65 70 75 80

Pro Pro Pro Gly Phe Ala His Gln Gln Asn His Asp Gly Asn Thr Glu 85 90 95

Tyr Gln Ser Pro Ile Leu Arg Gly Thr Phe Cys Ile Phe Phe Ser Ser 100 105 110

Asn Thr Asn Lys Pro Asn Tyr Pro Gly Lys Ala Val Gln Asn Leu Lys 115

Glu Tyr Ala His Ala Ile Thr Leu Arg Ser Gly Lys Ala Leu Pro Thr 130 135 140

Arg Glu Glu Pro Lys Thr Val Thr Glu Asp Ser Glu Asp Gln Asp Gly
145 150 155 160

Glu Asp Leu Ser Leu Glu Gln Pro Leu Asp Xaa Ser Leu Glu Xaa Pro
165 170 175

Leu Asp Gln Pro Leu Glu Gln Pro Leu Asp Arg Ala Thr Gln Pro Lleu

Leu Asp Gln Pro Leu Glu Gln Pro Leu Asp Arg Ala Thr Gln Pro Ile 180 185 190

Phe Pro Ala Val Ser Ser Thr 195

- (2) INFORMATION FOR SEQ ID NO:1175:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 126 amino acids

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..126
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597173
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1175:
- Trp Gly Arg Val Thr Ser Trp Lys Lys Ser Ala Thr Ser Thr Thr 1 5 10 15
- Arg Val Ala Thr Lys Asp Thr Thr Thr Ser Lys Pro Thr Ile Pro Thr 20 25 30
- Ser Pro Thr Val Ala Pro Ala Leu Leu Ile Leu Arg Ile Arg Cys Ile 35 40 45
- Pro Cys Asn Asn Asn Lys Val Arg Thr Asn Leu Leu Phe Ser Thr Thr 50 60
- Lys Val Ser Phe Leu Ser Ser Asn Xaa Trp Gly Thr Thr Ser Arg His 65 70 75 80
- His His Leu Asp Leu His Ile Ser Lys Thr Met Met Glu Thr Leu Asn 85 90 95
- Thr Lys Val Arg Tyr Leu Glu Gly His Ser Ala Ser Ser Ser Ala Pro 100 105 110
- Thr Gln Thr Ser Gln Thr Thr Gln Ala Lys Gln Phe Arg Ile 115 120 125
- (2) INFORMATION FOR SEQ ID NO:1176:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1026 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..1026
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597192
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1176:
- gaagaacccc gtaacgttta tctcgggttg tgtrctgatg ggttcaatcc atttggcatg 60 tetegtaate attegttgtg geetgtgate ctaacteett ataatttace ceetggtatg 120 tgcatgaata cagagtactt atttcttacc attctgaatt ctggtccaaa tcatcctcga 180 gctagtcttg atgtcttcct ccaacctctt attgaggagt taaaagagtt gtggtctact 240 ggagtcgatg cgtacgatgt ttcattgagt cagaatttta atctaaaagc agtactacta 300 tggacgatta gcgactttcc ggcgtatagc atgttatcag gatggactac tcagttttgt 360 aaatgtagta tgttcgaaga ttatctttct gcgaaatatc catgcttgcc cgaaaaagaa 420 ctctacgcga gaagagcgaa agaatatcat ctatgggtta aagaatatgt aacatactgg 480 aacactactt ctccatttcc tacttgggtt caagagattg tgcaaggacc tttgaacaag 540 gttaaaactt ggccaatgta tttcacaaga ggatatttgt ttcatacgca aaatcacggc 600 getggacgaa agacatgtaa etatggggte tgtgttaaag gtggaaatta egetgattea 660 tetgatgaag eggattteta eggtaeetta aetgataata tagaaetega gtatgaggat 720 caagttigtt atatteegta tecatataeg aagaaaceaa agaacattig geteaatgtt 780 ctaaaagtca atccgagggg aaatatttct ggagaatatg aaaacaatga tccaactctt 840 ttgcaaacag aaaatgatga tgatgttttg ctcactacaa ttgaagatct tgttcttgaa 900 actocggtag ccaacttaaa cccaataatc cttgattacg atgtcgggga tgctgaacca 960 gaagacgaat toogatgtaa tttatogtoo toggacgaag atgaagtaga agatgaagat 1020
- (2) INFORMATION FOR SEQ ID NO:1177:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 341 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..341
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597193
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1177: Glu Glu Pro Arg Asn Val Tyr Leu Gly Leu Cys Xaa Asp Gly Phe Asn 10 5 Pro Phe Gly Met Ser Arg Asn His Ser Leu Trp Pro Val Ile Leu Thr 25 Pro Tyr Asn Leu Pro Pro Gly Met Cys Met Asn Thr Glu Tyr Leu Phe 40 Leu Thr Ile Leu Asn Ser Gly Pro Asn His Pro Arg Ala Ser Leu Asp 55 Val Phe Leu Gln Pro Leu Ile Glu Glu Leu Lys Glu Leu Trp Ser Thr 75 70 Gly Val Asp Ala Tyr Asp Val Ser Leu Ser Gln Asn Phe Asn Leu Lys 90 85 Ala Val Leu Leu Trp Thr Ile Ser Asp Phe Pro Ala Tyr Ser Met Leu 105 Ser Gly Trp Thr Thr Gln Phe Cys Lys Cys Ser Met Phe Glu Asp Tyr 125 120 Leu Ser Ala Lys Tyr Pro Cys Leu Pro Glu Lys Glu Leu Tyr Ala Arg 135 140 Arg Ala Lys Glu Tyr His Leu Trp Val Lys Glu Tyr Val Thr Tyr Trp 155 150 Asn Thr Thr Ser Pro Phe Pro Thr Trp Val Gln Glu Ile Val Gln Gly 170 165 Pro Leu Asn Lys Val Lys Thr Trp Pro Met Tyr Phe Thr Arg Gly Tyr 185 180 Leu Phe His Thr Gln Asn His Gly Ala Gly Arg Lys Thr Cys Asn Tyr 200 205 Gly Val Cys Val Lys Gly Gly Asn Tyr Ala Asp Ser Ser Asp Glu Ala 215 220 Asp Phe Tyr Gly Thr Leu Thr Asp Asn Ile Glu Leu Glu Tyr Glu Asp 230 235 Gln Val Cys Tyr Ile Pro Tyr Pro Tyr Thr Lys Lys Pro Lys Asn Ile 245 250 Trp Leu Asn Val Leu Lys Val Asn Pro Arg Gly Asn Ile Ser Gly Glu 265 Tyr Glu Asn Asn Asp Pro Thr Leu Leu Gln Thr Glu Asn Asp Asp Asp 280 285 Val Leu Leu Thr Thr Ile Glu Asp Leu Val Leu Glu Thr Pro Val Ala 300 295 Asn Leu Asn Pro Ile Ile Leu Asp Tyr Asp Val Gly Asp Ala Glu Pro

325 Glu Asp Glu Asp Val 340

- (2) INFORMATION FOR SEQ ID NO:1178:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 322 amino acids

310 315

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..322
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597194

Glu Asp Glu Phe Arg Cys Asn Leu Ser Ser Asp Glu Asp Glu Val

Attorney Docket Na 750-1237P

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1178: Met Ser Arg Asn His Ser Leu Trp Pro Val Ile Leu Thr Pro Tyr Asn 10 Leu Pro Pro Gly Met Cys Met Asn Thr Glu Tyr Leu Phe Leu Thr Ile 20 25 Leu Asn Ser Gly Pro Asn His Pro Arg Ala Ser Leu Asp Val Phe Leu 45 40 Gln Pro Leu Ile Glu Glu Leu Lys Glu Leu Trp Ser Thr Gly Val Asp 55 Ala Tyr Asp Val Ser Leu Ser Gln Asn Phe Asn Leu Lys Ala Val Leu 75 Leu Trp Thr Ile Ser Asp Phe Pro Ala Tyr Ser Met Leu Ser Gly Trp 90 85 Thr Thr Gln Phe Cys Lys Cys Ser Met Phe Glu Asp Tyr Leu Ser Ala 105 100 Lys Tyr Pro Cys Leu Pro Glu Lys Glu Leu Tyr Ala Arg Arg Ala Lys 125 120 Glu Tyr His Leu Trp Val Lys Glu Tyr Val Thr Tyr Trp Asn Thr Thr 140 135 Ser Pro Phe Pro Thr Trp Val Gln Glu Ile Val Gln Gly Pro Leu Asn 155 150 Lys Val Lys Thr Trp Pro Met Tyr Phe Thr Arg Gly Tyr Leu Phe His 170 165 Thr Gln Asn His Gly Ala Gly Arg Lys Thr Cys Asn Tyr Gly Val Cys 185 180 Val Lys Gly Gly Asn Tyr Ala Asp Ser Ser Asp Glu Ala Asp Phe Tyr 205 200 Gly Thr Leu Thr Asp Asn Ile Glu Leu Glu Tyr Glu Asp Gln Val Cys 220 215 Tyr Ile Pro Tyr Pro Tyr Thr Lys Lys Pro Lys Asn Ile Trp Leu Asn 235 230 Val Leu Lys Val Asn Pro Arg Gly Asn Ile Ser Gly Glu Tyr Glu Asn 250 255 245 Asn Asp Pro Thr Leu Leu Gln Thr Glu Asn Asp Asp Val Leu Leu 270 265 260 Thr Thr Ile Glu Asp Leu Val Leu Glu Thr Pro Val Ala Asn Leu Asn 285 280 Pro Ile Ile Leu Asp Tyr Asp Val Gly Asp Ala Glu Pro Glu Asp Glu 300 295 Phe Arg Cys Asn Leu Ser Ser Ser Asp Glu Asp Glu Val Glu Asp Glu 315 310 305 Asp Val

- (2) INFORMATION FOR SEQ ID NO:1179:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 302 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..302
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597195
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1179:
 Met Cys Met Asn Thr Glu Tyr Leu Phe Leu Thr Ile Leu Asn Ser Gly
 10 15
- Pro Asn His Pro Arg Ala Ser Leu Asp Val Phe Leu Gln Pro Leu Ile
 20 25 30
- Glu Glu Leu Lys Glu Leu Trp Ser Thr Gly Val Asp Ala Tyr Asp Val 35

Ser Leu Ser Gln Asn Phe Asn Leu Lys Ala Val Leu Leu Trp Thr Ile 55 Ser Asp Phe Pro Ala Tyr Ser Met Leu Ser Gly Trp Thr Thr Gln Phe 75 70 Cys Lys Cys Ser Met Phe Glu Asp Tyr Leu Ser Ala Lys Tyr Pro Cys 90 85 Leu Pro Glu Lys Glu Leu Tyr Ala Arg Arg Ala Lys Glu Tyr His Leu 105 Trp Val Lys Glu Tyr Val Thr Tyr Trp Asn Thr Thr Ser Pro Phe Pro 125 120 115 Thr Trp Val Gln Glu Ile Val Gln Gly Pro Leu Asn Lys Val Lys Thr 140 135 Trp Pro Met Tyr Phe Thr Arg Gly Tyr Leu Phe His Thr Gln Asn His 155 150 Gly Ala Gly Arg Lys Thr Cys Asn Tyr Gly Val Cys Val Lys Gly Gly 170 165 Asn Tyr Ala Asp Ser Ser Asp Glu Ala Asp Phe Tyr Gly Thr Leu Thr 185 Asp Asn Ile Glu Leu Glu Tyr Glu Asp Gln Val Cys Tyr Ile Pro Tyr 200 195 Pro Tyr Thr Lys Lys Pro Lys Asn Ile Trp Leu Asn Val Leu Lys Val 220 215 Asn Pro Arg Gly Asn Ile Ser Gly Glu Tyr Glu Asn Asn Asp Pro Thr 235 230 Leu Leu Gln Thr Glu Asn Asp Asp Asp Val Leu Leu Thr Thr Ile Glu 250 245 Asp Leu Val Leu Glu Thr Pro Val Ala Asn Leu Asn Pro Ile Ile Leu 265 Asp Tyr Asp Val Gly Asp Ala Glu Pro Glu Asp Glu Phe Arg Cys Asn 275 280 Leu Ser Ser Ser Asp Glu Asp Glu Val Glu Asp Glu Asp Val 295

- (2) INFORMATION FOR SEQ ID NO:1180:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 240 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..240
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597208
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1180:

tccagtgagg tgtctcctgt tgtcttgatg ggcaatcggt taacacagtt gcttgatgcg 60 atggagctaa gtcggcaaac catgaagacc gtgaagcaaa acctttggtg ggcgttcgga 120 tacaacattg taggaatccc aattgcggct ggagtgttgc taccattgac tggaaccatg ttgactccat caatggcggg ggctctcatg ggntgtaagc tctctcggtg tcatgactaa 240

- (2) INFORMATION FOR SEQ ID NO:1181:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 79 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..79
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597209
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1181:

Client Docket No. 55146.003 Ser Ser Glu Val Ser Pro Val Val Leu Met Gly Asn Arg Leu Thr Gln 10 5 Leu Leu Asp Ala Met Glu Leu Ser Arg Gln Thr Met Lys Thr Val Lys 25 20 Gln Asn Leu Trp Trp Ala Phe Gly Tyr Asn Ile Val Gly Ile Pro Ile 40 Ala Ala Gly Val Leu Leu Pro Leu Thr Gly Thr Met Leu Thr Pro Ser 55 Met Ala Gly Ala Leu Met Xaa Cys Lys Leu Ser Arg Cys His Asp 70 (2) INFORMATION FOR SEQ ID NO:1182: (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 70 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..70
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597210
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1182:

Met Gly Asn Arg Leu Thr Gln Leu Leu Asp Ala Met Glu Leu Ser Arg 10 5

Gln Thr Met Lys Thr Val Lys Gln Asn Leu Trp Trp Ala Phe Gly Tyr 20 25

Asn Ile Val Gly Ile Pro Ile Ala Ala Gly Val Leu Leu Pro Leu Thr 45 4.0

Gly Thr Met Leu Thr Pro Ser Met Ala Gly Ala Leu Met Xaa Cys Lys 55

Leu Ser Arg Cys His Asp 70

- (2) INFORMATION FOR SEQ ID NO:1183:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 59 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..59
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597211
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1183:

Met Glu Leu Ser Arg Gln Thr Met Lys Thr Val Lys Gln Asn Leu Trp 1.0 5

Trp Ala Phe Gly Tyr Asn Ile Val Gly Ile Pro Ile Ala Ala Gly Val 25

Leu Leu Pro Leu Thr Gly Thr Met Leu Thr Pro Ser Met Ala Gly Ala 40

Leu Met Xaa Cys Lys Leu Ser Arg Cys His Asp 55

- (2) INFORMATION FOR SEQ ID NO:1184:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 982 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -



(B) LOCATION: 1..982
(D) OTHER INFORMATION: / Ceres Seq. ID 1597224

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1184: 60 aacacaattt gatacgcgtc agtttgacca gaggctgaat gaagttctcg atggacagga 120 tgagttcttc acctcatatg atgaggtcca tgagagcttt gatgccatgg gtttgcaaga 180 qaatcttctt aggggtatct atgcttacgg tttcgaaaag ccttctgcta ttcagcaaag 240 aggaattgta cccttttgca agggtcttga tgtgatccag caggcacagt ctggtactgg 300 aaaaaccgcc actttctgct ctggtgtctt gcagcagctt gactatgccc ttctccagtg 360 ccaggetete gttttggete ccaecagaga gettgeteag cagattgaga aggteatgeg tgcccttggt gactaccaag gtgtcaaggt tcatgcctgt gttggtggaa ccagtgtccg 420 tnaggateag egeattetee aggetggtgt teatgttgte gttggaacte etggtegtgt 480 540 ttttqacatq cttaqaaqac aatctctccg ccctgactgc atcaagatgt ttgtccttga 600 tgaagetgat gaaatgetet eeegtggttt caaggateag atetatgaea tatteeaget tctcccacca aagattcagg ttggagtgtt ctctgcgaca atgcctccgg aagctttgga 660 gatcacaagg aaattcatga gcaaaccagt gagaatcttg gtgaaacgtg atgagctcac 720 acttgaaggn tatcaagcaa ttctacgtga ancgtggaga aagaagactg gaangcttga 780 gactctctgc gatctctacg nagactctag ccatcactca gagtgtcatc tttgtcaaca 840 ctcgtcgcaa ggtcganctg gctncacaga caaaatgaga agccgtgacc acacagtctc 900 agennnetca tggagacatg gaccaaaaca ccagagacat catcatgaga gagttcaggt 960 cntggctcgt ctcgtgttct aa

- (2) INFORMATION FOR SEQ ID NO:1185:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 326 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..326
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597225
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1185: Thr Gln Phe Asp Thr Arg Gln Phe Asp Gln Arg Leu Asn Glu Val Leu
- 10 15
 Asp Gly Gln Asp Glu Phe Phe Thr Ser Tyr Asp Glu Val His Glu Ser 20 25 30
 Phe Asp Ala Met Gly Leu Gln Glu Asn Leu Leu Arg Gly Ile Tyr Ala 35
 Tyr Gly Phe Glu Lys Pro Ser Ala Ile Gln Gln Arg Gly Ile Val Pro 50
- 50 55 60

 Phe Cys Lys Gly Leu Asp Val Ile Gln Gln Ala Gln Ser Gly Thr Gly
 65 70 75 80
- Lys Thr Ala Thr Phe Cys Ser Gly Val Leu Gln Gln Leu Asp Tyr Ala
 85 90 95
- Leu Leu Gln Cys Gln Ala Leu Val Leu Ala Pro Thr Arg Glu Leu Ala 100 105 110
- Gln Gln Ile Glu Lys Val Met Arg Ala Leu Gly Asp Tyr Gln Gly Val 115 120 125
- Lys Val His Ala Cys Val Gly Gly Thr Ser Val Arg Xaa Asp Gln Arg
- Ile Leu Gln Ala Gly Val His Val Val Gly Thr Pro Gly Arg Val
 145 150 155 160
- Phe Asp Met Leu Arg Arg Gln Ser Leu Arg Pro Asp Cys Ile Lys Met

 165 170 175
- Phe Val Leu Asp Glu Ala Asp Glu Met Leu Ser Arg Gly Phe Lys Asp 180 185 190
- Gln Ile Tyr Asp Ile Phe Gln Leu Leu Pro Pro Lys Ile Gln Val Gly
 195 200 205
- Val Phe Ser Ala Thr Met Pro Pro Glu Ala Leu Glu Ile Thr Arg Lys 210 215 220

 Phe Met Ser Lys Pro Val Arg Ile Leu Val Lys Arg Asp Glu Leu Thr

230 235 225 Leu Glu Xaa Tyr Gln Ala Ile Leu Arg Glu Xaa Trp Arg Lys Lys Thr 245 250 Gly Xaa Leu Glu Thr Leu Cys Asp Leu Tyr Xaa Asp Ser Ser His His 265 Ser Glu Cys His Leu Cys Gln His Ser Ser Gln Gly Arg Xaa Gly Xaa 280 285 Thr Asp Lys Met Arg Ser Arg Asp His Thr Val Ser Xaa Xaa Ser Trp 295 Arg His Gly Pro Lys His Gln Arg His His His Glu Arg Val Gln Val 315 310 Xaa Ala Arg Leu Val Phe

325

- (2) INFORMATION FOR SEQ ID NO:1186:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 291 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..291
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597226
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1186: Met Gly Leu Gln Glu Asn Leu Leu Arg Gly Ile Tyr Ala Tyr Gly Phe 10 5 Glu Lys Pro Ser Ala Ile Gln Gln Arg Gly Ile Val Pro Phe Cys Lys 20 25 Gly Leu Asp Val Ile Gln Gln Ala Gln Ser Gly Thr Gly Lys Thr Ala 40 Thr Phe Cys Ser Gly Val Leu Gln Gln Leu Asp Tyr Ala Leu Leu Gln 5.5 Cys Gln Ala Leu Val Leu Ala Pro Thr Arg Glu Leu Ala Gln Gln Ile 70 75 Glu Lys Val Met Arg Ala Leu Gly Asp Tyr Gln Gly Val Lys Val His 90 85 Ala Cys Val Gly Gly Thr Ser Val Arg Xaa Asp Gln Arg Ile Leu Gln 105 110 100 Ala Gly Val His Val Val Val Gly Thr Pro Gly Arg Val Phe Asp Met 120 125 Leu Arg Arg Gln Ser Leu Arg Pro Asp Cys Ile Lys Met Phe Val Leu 140 135 Asp Glu Ala Asp Glu Met Leu Ser Arg Gly Phe Lys Asp Gln Ile Tyr 150 155 Asp Ile Phe Gln Leu Leu Pro Pro Lys Ile Gln Val Gly Val Phe Ser 170 175 165 Ala Thr Met Pro Pro Glu Ala Leu Glu Ile Thr Arg Lys Phe Met Ser 185 190 180 Lys Pro Val Arg Ile Leu Val Lys Arg Asp Glu Leu Thr Leu Glu Xaa 200 205 Tyr Gln Ala Ile Leu Arg Glu Xaa Trp Arg Lys Lys Thr Gly Xaa Leu 215 220 Glu Thr Leu Cys Asp Leu Tyr Xaa Asp Ser Ser His His Ser Glu Cys 235 230 His Leu Cys Gln His Ser Ser Gln Gly Arg Xaa Gly Xaa Thr Asp Lys 250 245 Met Arg Ser Arg Asp His Thr Val Ser Xaa Xaa Ser Trp Arg His Gly 265 Pro Lys His Gln Arg His His His Glu Arg Val Gln Val Xaa Ala Arg 280 285

Leu Val Phe 290

- (2) INFORMATION FOR SEQ ID NO:1187:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 208 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..208
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597227
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1187: Met Arg Ala Leu Gly Asp Tyr Gln Gly Val Lys Val His Ala Cys Val 10 5 Gly Gly Thr Ser Val Arg Xaa Asp Gln Arg Ile Leu Gln Ala Gly Val 2.0 25 His Val Val Gly Thr Pro Gly Arg Val Phe Asp Met Leu Arg Arg 40 Gln Ser Leu Arg Pro Asp Cys Ile Lys Met Phe Val Leu Asp Glu Ala 60 55 Asp Glu Met Leu Ser Arg Gly Phe Lys Asp Gln Ile Tyr Asp Ile Phe 75 70 Gln Leu Leu Pro Pro Lys Ile Gln Val Gly Val Phe Ser Ala Thr Met 90 85 Pro Pro Glu Ala Leu Glu Ile Thr Arg Lys Phe Met Ser Lys Pro Val 110 105 Arg Ile Leu Val Lys Arg Asp Glu Leu Thr Leu Glu Xaa Tyr Gln Ala 120 125 Ile Leu Arg Glu Xaa Trp Arg Lys Lys Thr Gly Xaa Leu Glu Thr Leu 140 135 Cys Asp Leu Tyr Xaa Asp Ser Ser His His Ser Glu Cys His Leu Cys 155 150 Gln His Ser Ser Gln Gly Arg Xaa Gly Xaa Thr Asp Lys Met Arg Ser 165 170 Arg Asp His Thr Val Ser Xaa Xaa Ser Trp Arg His Gly Pro Lys His 190 185

Gln Arg His His Glu Arg Val Gln Val Xaa Ala Arg Leu Val Phe

200

205

- (2) INFORMATION FOR SEQ ID NO:1188:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 342 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..342
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597228
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1188:

 atgaagttg acatcagaga agcttcaggg aaatcaacaa caggagttca gaacatnaat 60

 attgatgaag nngcagaagc tggaacccca ccaagagtta actatacttc tcaacttagc 120

 aagctgaaga ggacttttga ccacaagaag agagcaacag aaaggctagc tcagactgga 180

 gatcctacaa aagatgactg ctctgtaaga gtcaagctta gtgaattaaa acaagctcac 240

 ttgggaggaa gtcccatgtc tatccaggnt tgtctcattg atcattataa agtcaatggt 300

 cgagtnaact caaaccaagt caagaaagat ctcagccgtt ga

(2) INFORMATION FOR SEQ ID NO:1189:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 113 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..113
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597229
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1189:

Met Lys Phe Asp Ile Arg Glu Ala Ser Gly Lys Ser Thr Thr Gly Val

Gln Asn Xaa Asn Ile Asp Glu Xaa Ala Glu Ala Gly Thr Pro Pro Arg
20 25 30 ...

Val Asn Tyr Thr Ser Gln Leu Ser Lys Leu Lys Arg Thr Phe Asp His 35

Lys Lys Arg Ala Thr Glu Arg Leu Ala Gln Thr Gly Asp Pro Thr Lys 50 60

Asp Asp Cys Ser Val Arg Val Lys Leu Ser Glu Leu Lys Gln Ala His 65 70 75 80
Leu Gly Gly Ser Pro Met Ser Ile Gln Xaa Cys Leu Ile Asp His Tyr

85 90 95

Lie Val Agn Cly Arg Van Agn Ser Agn Gln Val Lvs Lvs Agn Leu Ser

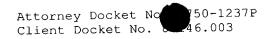
Lys Val Asn Gly Arg Xaa Asn Ser Asn Gln Val Lys Lys Asp Leu Ser 100 105 110

Arg

- (2) INFORMATION FOR SEQ ID NO:1190:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1032 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..1032
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597257
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1190:

atngentgtn atcattnetg ettngteatn cactngcetg etatngeeat ttetgetttt 60 agtituatt cattucgeti attgcatact aggacigtta gaaccaacca acttictact 120 180 tggcttgact tagagctttc tgatcatatc tcatctgttt gcgtcacact gaattggatt gacatectaa aatactacaa egacaggata gtacettgca acctgettta cagacgcate 240 tetgattegt etgttatage cenggetget atetgttgea tgeatacaeg gteagaagga 300 aatcagaacc tectatteaa egataacate gacegtattg etegecaact aagagaacag 360 acagaaaccg acacaatggc tgacgttgta gatgagcaag agcaacctac caacattggt 420 getggtgact teceteacaa ecacaaccag egteatggaa tigtteeace tecagtacag 480 540 aacaacaact ttgagatcaa aagcggtctc attgctatgg ttcaagggaa caagtttcat ggcctgccaa tggaggatcc gctagatcat ctngacgagt tngaaagnet ctgtgncctn 600 actaaaatca atggagttag tgaagatggt ttnnnanttc gcttgtttcc attctcactt 660 ggagataaag cccatctgtg ggaaaagacg ctaccccaga attcaatcac aacctgggat 720 gactgcaaaa aggccttctt ggcaaaattc ttttccaact ccagaactgc aagactccgg 780 aatgagatat coggatttac toagaagcaa aatgaaagct totgtgaagc ttgggagcgc 840 tttaagggtt atcaaaccaa atgccctcat cacggattta agcangctnc tctnctcagc 900 acactetata gagnegtetn geetaagata aggatactge ttgacacege ttcaaatggg 960 aattttttga agaaggatgt tgaagaagga tggnagctag tngaaaanct ttgctcagtc 1020 gnatggcaat na

- (2) INFORMATION FOR SEQ ID NO:1191:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 343 amino acids
 - (B) TYPE: amino acid



- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..343
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597258
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1191:
- Xaa Xaa Cys Xaa His Xaa Cys Xaa Val Xaa His Xaa Pro Ala Xaa Ala 10 5
- Ile Ser Ala Phe Ser Phe Ile Ser Phe Arg Leu Leu His Thr Arg Thr 25
- Val Arg Thr Asn Gln Leu Ser Thr Trp Leu Asp Leu Glu Leu Ser Asp 45 40
- His Ile Ser Ser Val Cys Val Thr Leu Asn Trp Ile Asp Ile Leu Lys 55
- Tyr Tyr Asn Asp Arg Ile Val Pro Cys Asn Leu Leu Tyr Arg Arg Ile 75 65 70
- Ser Asp Ser Ser Val Ile Ala Xaa Ala Ala Ile Cys Cys Met His Thr 90 85
- Arg Ser Glu Gly Asn Gln Asn Leu Leu Phe Asn Asp Asn Ile Asp Arg 105
- Ile Ala Arg Gln Leu Arg Glu Gln Thr Glu Thr Asp Thr Met Ala Asp
- 125 120 Val Val Asp Glu Gln Glu Gln Pro Thr Asn Ile Gly Ala Gly Asp Phe 140
- 130 135 Pro His Asn His Asn Gln Arg His Gly Ile Val Pro Pro Pro Val Gln 155
- 150 Asn Asn Asn Phe Glu Ile Lys Ser Gly Leu Ile Ala Met Val Gln Gly 170
- 165 Asn Lys Phe His Gly Leu Pro Met Glu Asp Pro Leu Asp His Xaa Asp 180 185 190
- Glu Xaa Glu Xaa Leu Cys Xaa Xaa Thr Lys Ile Asn Gly Val Ser Glu 200 205
- Asp Gly Xaa Xaa Xaa Arg Leu Phe Pro Phe Ser Leu Gly Asp Lys Ala 210 215 220
- His Leu Trp Glu Lys Thr Leu Pro Gln Asn Ser Ile Thr Trp Asp 230 235
- Asp Cys Lys Lys Ala Phe Leu Ala Lys Phe Phe Ser Asn Ser Arg Thr 250 245
- Ala Arg Leu Arg Asn Glu Ile Ser Gly Phe Thr Gln Lys Gln Asn Glu 260 265 Ser Phe Cys Glu Ala Trp Glu Arg Phe Lys Gly Tyr Gln Thr Lys Cys
- 285 280
- Pro His His Gly Phe Lys Xaa Ala Xaa Xaa Leu Ser Thr Leu Tyr Arg 295 300
- Xaa Val Xaa Pro Lys Ile Arg Ile Leu Leu Asp Thr Ala Ser Asn Gly 310 315 320
- Asn Phe Leu Lys Lys Asp Val Glu Glu Gly Trp Xaa Leu Xaa Glu Xaa 325 330

Leu Cys Ser Val Xaa Trp Gln

- (2) INFORMATION FOR SEQ ID NO:1192:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 250 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide

(B) LOCATION: 1..250 (D) OTHER INFORMATION: / Ceres Seq. ID 1597259 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1192: Met His Thr Arg Ser Glu Gly Asn Gln Asn Leu Leu Phe Asn Asp Asn 10 Ile Asp Arg Ile Ala Arg Gln Leu Arg Glu Gln Thr Glu Thr Asp Thr Met Ala Asp Val Val Asp Glu Gln Glu Gln Pro Thr Asn Ile Gly Ala 40 Gly Asp Phe Pro His Asn His Asn Gln Arg His Gly Ile Val Pro Pro 55 Pro Val Gln Asn Asn Asn Phe Glu Ile Lys Ser Gly Leu Ile Ala Met 70 Val Gln Gly Asn Lys Phe His Gly Leu Pro Met Glu Asp Pro Leu Asp 8.5 His Xaa Asp Glu Xaa Glu Xaa Leu Cys Xaa Xaa Thr Lys Ile Asn Gly 105 Val Ser Glu Asp Gly Xaa Xaa Xaa Arg Leu Phe Pro Phe Ser Leu Gly 120 Asp Lys Ala His Leu Trp Glu Lys Thr Leu Pro Gln Asn Ser Ile Thr 135 140 Thr Trp Asp Asp Cys Lys Lys Ala Phe Leu Ala Lys Phe Phe Ser Asn 150 155 Ser Arg Thr Ala Arg Leu Arg Asn Glu Ile Ser Gly Phe Thr Gln Lys 165 170 Gln Asn Glu Ser Phe Cys Glu Ala Trp Glu Arg Phe Lys Gly Tyr Gln 185 190 Thr Lys Cys Pro His His Gly Phe Lys Xaa Ala Xaa Xaa Leu Ser Thr 200 205

Leu Tyr Arg Xaa Val Xaa Pro Lys Ile Arg Ile Leu Leu Asp Thr Ala

Ser Asn Gly Asn Phe Leu Lys Lys Asp Val Glu Glu Gly Trp Xaa Leu

220

- 245 (2) INFORMATION FOR SEQ ID NO:1193:
 - (i) SEQUENCE CHARACTERISTICS:

Xaa Glu Xaa Leu Cys Ser Val Xaa Trp Gln

(A) LENGTH: 218 amino acids

215

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..218
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597260
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1193:
- Met Ala Asp Val Val Asp Glu Gln Glu Gln Pro Thr Asn Ile Gly Ala 10 Gly Asp Phe Pro His Asn His Asn Gln Arg His Gly Ile Val Pro Pro
- 25
- Pro Val Gln Asn Asn Asn Phe Glu Ile Lys Ser Gly Leu Ile Ala Met 40
- Val Gln Gly Asn Lys Phe His Gly Leu Pro Met Glu Asp Pro Leu Asp 55
- His Xaa Asp Glu Xaa Glu Xaa Leu Cys Xaa Xaa Thr Lys Ile Asn Gly 75 70
- Val Ser Glu Asp Gly Xaa Xaa Xaa Arg Leu Phe Pro Phe Ser Leu Gly 90 85
- Asp Lys Ala His Leu Trp Glu Lys Thr Leu Pro Gln Asn Ser Ile Thr 105 100

Thr Trp Asp Asp Cys Lys Lys Ala Phe Leu Ala Lys Phe Phe Ser Asn
115
120
125
127
128
129
129
125

Ser Arg Thr Ala Arg Leu Arg Asn Glu Ile Ser Gly Phe Thr Gln Lys
130
135
140

Gln Asn Glu Ser Phe Cys Glu Ala Trp Glu Arg Phe Lys Gly Tyr Gln 145 150 155 160

Thr Lys Cys Pro His His Gly Phe Lys Xaa Ala Xaa Xaa Leu Ser Thr 165 170 175

Leu Tyr Arg Xaa Val Xaa Pro Lys Ile Arg Ile Leu Leu Asp Thr Ala 180 185 190

Ser Asn Gly Asn Phe Leu Lys Lys Asp Val Glu Glu Gly Trp Xaa Leu 195 200 205

Xaa Glu Xaa Leu Cys Ser Val Xaa Trp Gln 210 215

- (2) INFORMATION FOR SEQ ID NO:1194:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 591 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..591
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597261
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1194: 60 atgtcaacaa gatcgccact cgcaaggatt caacctacaa gaacgttggt tggacttccc 120 ngtggctaca aaggtcgcga ctttnctcat cgcngaggac ctagttccat ctctcccngc aattctcaag attctttgaa caaagatgtt tgctttcacc ntggagtgaa catactcgca gctacttctc caaatgatac attttatgac ggaggatttc ctatgaaatc aggaacatca 240 atgtcgactc cttttgttgc aggaattgta gcactcctta agtctttgca ccctcattgg teteeggeeg ceattagate tgetattgtt actacagett ggagaacaga tecateegag aaagccgcga aacccggtct tgtttacgat atgggagtta acgattatgt tctntatttg 420 480 tgctccgttg gttacacaga ttcatccatt actagacttg tccgcaagaa gacggtttgt gcaaacccta aaccttcagt tcttgacctc aaattgcctt caatcacaat cccaaacctt 540 gcaaaannga agttattatc actagaaccg ttaccaacgt tggaccagta g
- (2) INFORMATION FOR SEQ ID NO:1195:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 196 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..196
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597262
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1195:
- Met Ser Thr Arg Ser Pro Leu Ala Arg Ile Gln Pro Thr Arg Thr Leu 1 5 10 15
- Val Gly Leu Pro Xaa Gly Tyr Lys Gly Arg Asp Phe Xaa His Arg Xaa 20 25 30
- Gly Pro Ser Ser Ile Ser Pro Xaa Asn Ser Gln Asp Ser Leu Asn Lys
- Asp Val Cys Phe His Xaa Gly Val Asn Ile Leu Ala Ala Thr Ser Pro
 50 60
- Asn Asp Thr Phe Tyr Asp Gly Gly Phe Pro Met Lys Ser Gly Thr Ser 65 70 75 80

 Met Ser Thr Pro Phe Val Ala Gly Ile Val Ala Leu Leu Lys Ser Leu
- 85 90 95
 His Pro His Trp Ser Pro Ala Ala Ile Arg Ser Ala Ile Val Thr Thr

105 100 Ala Trp Arg Thr Asp Pro Ser Glu Lys Ala Ala Lys Pro Gly Leu Val 125 120 Tyr Asp Met Gly Val Asn Asp Tyr Val Xaa Tyr Leu Cys Ser Val Gly

140 135 Tyr Thr Asp Ser Ser Ile Thr Arg Leu Val Arg Lys Lys Thr Val Cys 155 150

Ala Asn Pro Lys Pro Ser Val Leu Asp Leu Lys Leu Pro Ser Ile Thr 170 165

Ile Pro Asn Leu Ala Lys Xaa Lys Leu Leu Ser Leu Glu Pro Leu Pro 185 180

Thr Leu Asp Gln 195

- (2) INFORMATION FOR SEQ ID NO:1196:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 122 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..122
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597264

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1196:

Met Lys Ser Gly Thr Ser Met Ser Thr Pro Phe Val Ala Gly Ile Val 10 Ala Leu Leu Lys Ser Leu His Pro His Trp Ser Pro Ala Ala Ile Arg

30 25 20

Ser Ala Ile Val Thr Thr Ala Trp Arg Thr Asp Pro Ser Glu Lys Ala 45 40

Ala Lys Pro Gly Leu Val Tyr Asp Met Gly Val Asn Asp Tyr Val Xaa 60 55

Tyr Leu Cys Ser Val Gly Tyr Thr Asp Ser Ser Ile Thr Arg Leu Val 75 70

Arg Lys Lys Thr Val Cys Ala Asn Pro Lys Pro Ser Val Leu Asp Leu 90 85

Lys Leu Pro Ser Ile Thr Ile Pro Asn Leu Ala Lys Xaa Lys Leu Leu 105 100

Ser Leu Glu Pro Leu Pro Thr Leu Asp Gln

- (2) INFORMATION FOR SEQ ID NO:1197:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 714 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..714
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597281
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1197:

tetgtetgte tegteactea ggateagate ttagetaaat tgagacaage teaaaatgag taagetteag agtgaggeeg ttegtgaage cateactact ateacaggga aateegagge 120 aaagaaacgt aactttgtcg agactattga gctccagatc ggtctgaaga actatgaccc 180 tcaaaaggac aagcgtttca gtggatctgt caagttacca catatccccc gtcctaaaat 240 gaagatetge atgeteggag atgeceagea tgttgaagag gaaaatteee aactettgtg 300 agccaccagg aatccttgga gtcaaaggtg aatgaaacaa aggcaacagt gaagttccag 360 ctgaagaagg ttctgtgcat gggagttgca gttggtaacc tttcaatgga agagaagcag 420 atctttcaga atgtgcagat gagcgtcaac ttcctcgtct cgctattgaa gaagaactgg 480 caaaatgtca ggtgtttgta cctcaagagc acaatgggac caccacaaag aatcttctga 540 gctttcacct acttaaaatt ggtggctata agacacttct tcttaccctt tcgtttttgt 600 ttcagacata tctgttaaac agtgtaagag ttgtgctgct actctcctct ttagcttttg gtttggataa tttgaaataa ctgcagtgat tttggttaat ggattttgta gctt

- (2) INFORMATION FOR SEQ ID NO:1198:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 81 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..81
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597282
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1198:

Met Ser Lys Leu Gln Ser Glu Ala Val Arg Glu Ala Ile Thr Thr Ile

Thr Gly Lys Ser Glu Ala Lys Lys Arg Asn Phe Val Glu Thr Ile Glu 20 25 30

Leu Gln Ile Gly Leu Lys Asn Tyr Asp Pro Gln Lys Asp Lys Arg Phe

Ser Gly Ser Val Lys Leu Pro His Ile Pro Arg Pro Lys Met Lys Ile
50 55 60

Cys Met Leu Gly Asp Ala Gln His Val Glu Glu Glu Asn Ser Gln Leu 65 70 75 80

Leu

- (2) INFORMATION FOR SEQ ID NO:1199:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 53 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..53
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597283
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1199:

Gln Asn Val Gln Met Ser Val Asn Phe Leu Val Ser Leu Leu Lys Lys 20 25 30

Asn Trp Gln Asn Val Arg Cys Leu Tyr Leu Lys Ser Thr Met Gly Pro 35 40 45

Pro Gln Arg Ile Phe

50

- (2) INFORMATION FOR SEQ ID NO:1200:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 44 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..44
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597284
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1200:

Met Glu Glu Lys Gln Ile Phe Gln Asn Val Gln Met Ser Val Asn Phe

1 5 10 15
Leu Val Ser Leu Leu Lys Lys Asn Trp Gln Asn Val Arg Cys Leu Tyr
20 25 30

Leu Lys Ser Thr Met Gly Pro Pro Gln Arg Ile Phe 35 40

- (2) INFORMATION FOR SEQ ID NO:1201:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 821 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..821
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597285
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1201: 60 attgagtctt tagggtctct ctgtgtttgc aaatctgacg ccgagtattt tgctaaagca gcagccagtc gggacagtga aatggcgact gaagatgtgc aagatcccag aatcgctaag 120 attgcctctt ccattagagt catccccgac ttccctaaac caggaatcat gtttcaggac 180 ataacgacgc ttcttctcga cactgaggcc tttaaggata ctattgcttt gtttgttgat 240 agatacaaag ataaaggcat atctgttgtt gcaggtgttg aagctagagg tttcattttt 300 ggccctccta ttgcgttggc tattggtgcc aaatttgttc ccatgaggac gcccaagaag 360 ctacctggga aggttatttc ggaggagtat tcgttggagt atggaccaga tacgattgag 420 atgcacgtag gtgcagtaga gcctggtgag cgtgctatta ttattgatgn cctcattgcc 480 acgggtggga ctctcgctgc tgcaatccga ctacttgaac gagtaggagt gnagattgtt 540 gaatgtgctt gcgtaattga gttaccagag cttaagggaa aggagaaact aggagagacg 600 tcgctatttg ttcttgtaaa gtcggctgct taacaagaaa ctggaagtga aggttattgg 660 720 atcgagtgtt gatgctattt ttcatgtatg gtgagacatt ttgcgtggga tttgatcctt gttgnttcaa cttttatcat anttggttca gnctagaaaa tggcatttga atgtcaggat 780 togattgcag tttcctattg ttccacttaa aatccccgat t
- (2) INFORMATION FOR SEQ ID NO:1202:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 210 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..210
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597286

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1202:

Phe Ala Lys Ala Ala Ala Ser Arg Asp Ser Glu Met Ala Thr Glu Asp 20 25 30

Val Gln Asp Pro Arg Ile Ala Lys Ile Ala Ser Ser Ile Arg Val Ile 35 40 45

Pro Asp Phe Pro Lys Pro Gly Ile Met Phe Gln Asp Ile Thr Thr Leu 50 55

Leu Leu Asp Thr Glu Ala Phe Lys Asp Thr Ile Ala Leu Phe Val Asp 65 70 75 80

Arg Tyr Lys Asp Lys Gly Ile Ser Val Val Ala Gly Val Glu Ala Arg 90 95

Gly Phe Ile Phe Gly Pro Pro Ile Ala Leu Ala Ile Gly Ala Lys Phe $100 \hspace{1cm} 105 \hspace{1cm} 110$

Val Pro Met Arg Thr Pro Lys Lys Leu Pro Gly Lys Val Ile Ser Glu 115 120 125

Glu Tyr Ser Leu Glu Tyr Gly Pro Asp Thr Ile Glu Met His Val Gly 130 135 140

Ala Val Glu Pro Gly Glu Arg Ala Ile Ile Ile Asp Xaa Leu Ile Ala 150 155 Thr Gly Gly Thr Leu Ala Ala Ile Arg Leu Leu Glu Arg Val Gly 165 170 175

Val Xaa Ile Val Glu Cys Ala Cys Val Ile Glu Leu Pro Glu Leu Lys 190 180 185

Gly Lys Glu Lys Leu Gly Glu Thr Ser Leu Phe Val Leu Val Lys Ser 205 200

Ala Ala 210

- (2) INFORMATION FOR SEQ ID NO:1203:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 183 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..183
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597287
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1203:

Met Ala Thr Glu Asp Val Gln Asp Pro Arg Ile Ala Lys Ile Ala Ser 10 5

Ser Ile Arg Val Ile Pro Asp Phe Pro Lys Pro Gly Ile Met Phe Gln 30 25 20

Asp Ile Thr Thr Leu Leu Leu Asp Thr Glu Ala Phe Lys Asp Thr Ile 4.0

Ala Leu Phe Val Asp Arg Tyr Lys Asp Lys Gly Ile Ser Val Val Ala 55

Gly Val Glu Ala Arg Gly Phe Ile Phe Gly Pro Pro Ile Ala Leu Ala 75 70

Ile Gly Ala Lys Phe Val Pro Met Arg Thr Pro Lys Lys Leu Pro Gly 90 Lys Val Ile Ser Glu Glu Tyr Ser Leu Glu Tyr Gly Pro Asp Thr Ile

105 Glu Met His Val Gly Ala Val Glu Pro Gly Glu Arg Ala Ile Ile Ile 125

120 Asp Xaa Leu Ile Ala Thr Gly Gly Thr Leu Ala Ala Ala Ile Arg Leu 140 135

Leu Glu Arg Val Gly Val Xaa Ile Val Glu Cys Ala Cys Val Ile Glu 150 155 160

Leu Pro Glu Leu Lys Gly Lys Glu Lys Leu Gly Glu Thr Ser Leu Phe 170 165

Val Leu Val Lys Ser Ala Ala 180

(2) INFORMATION FOR SEQ ID NO:1204:

- - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 154 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..154
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597288

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1204:

Met Phe Gln Asp Ile Thr Thr Leu Leu Leu Asp Thr Glu Ala Phe Lys 5 10 Asp Thr Ile Ala Leu Phe Val Asp Arg Tyr Lys Asp Lys Gly Ile Ser

20 25 Val Val Ala Gly Val Glu Ala Arg Gly Phe Ile Phe Gly Pro Pro Ile 4.5 40 Ala Leu Ala Ile Gly Ala Lys Phe Val Pro Met Arg Thr Pro Lys Lys 55 Leu Pro Gly Lys Val Ile Ser Glu Glu Tyr Ser Leu Glu Tyr Gly Pro 75 70 Asp Thr Ile Glu Met His Val Gly Ala Val Glu Pro Gly Glu Arg Ala 85 Ile Ile Ile Asp Xaa Leu Ile Ala Thr Gly Gly Thr Leu Ala Ala Ala 105 100 Ile Arg Leu Leu Glu Arg Val Gly Val Xaa Ile Val Glu Cys Ala Cys 120 Val Ile Glu Leu Pro Glu Leu Lys Gly Lys Glu Lys Leu Gly Glu Thr 135 Ser Leu Phe Val Leu Val Lys Ser Ala Ala 150 (2) INFORMATION FOR SEQ ID NO:1205: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 365 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: -(B) LOCATION: 1..365 (D) OTHER INFORMATION: / Ceres Seq. ID 1597313 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1205: taagcttcag agtgaggccg ttcgtgaagc catcactact atcacaggga aatccgaggc 60 aaagaaacgt aactttgtcg agactattga gctccagatc ggtctgaaga actatgaccc 120 tcaaaaggac aagcgtttca gtggatctgt caagttacca catatccccc gtcctaaaat 180 gaagatetge atgeteggag atgeceagea tgttgaagag aacatggatg ttgagtetet 240 aaaaaagctt aacaagaaca agaaactcgt caagaagctt gcaaagaaat accatgcttt 300 cttggcctct gagtctgtca ttaagcagat tcctcgtctt cttggtcctg gtcttaacaa 360 ggcag (2) INFORMATION FOR SEQ ID NO:1206: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 121 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..121 (D) OTHER INFORMATION: / Ceres Seq. ID 1597314 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1206: Lys Leu Gln Ser Glu Ala Val Arg Glu Ala Ile Thr Thr Ile Thr Gly 5 10 Lys Ser Glu Ala Lys Lys Arg Asn Phe Val Glu Thr Ile Glu Leu Gln 25 Ile Gly Leu Lys Asn Tyr Asp Pro Gln Lys Asp Lys Arg Phe Ser Gly 40 Ser Val Lys Leu Pro His Ile Pro Arg Pro Lys Met Lys Ile Cys Met 60 55 Leu Gly Asp Ala Gln His Val Glu Glu Asn Met Asp Val Glu Ser Leu

75

90

Lys Lys Leu Asn Lys Asn Lys Lys Leu Val Lys Lys Leu Ala Lys Lys

Tyr His Ala Phe Leu Ala Ser Glu Ser Val Ile Lys Gln Ile Pro Arg

70

100 105 110

Leu Leu Gly Pro Gly Leu Asn Lys Ala 115 120

- (2) INFORMATION FOR SEQ ID NO:1207:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 62 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..62
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597315
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1207:

Asp Val Glu Ser Leu Lys Lys Leu Asn Lys Asn Lys Lys Leu Val Lys 20 25 30

Lys Leu Ala Lys Lys Tyr His Ala Phe Leu Ala Ser Glu Ser Val Ile 35 40 45

Lys Gln Ile Pro Arg Leu Leu Gly Pro Gly Leu Asn Lys Ala 50 55 60

- (2) INFORMATION FOR SEQ ID NO:1208:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1036 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..1036
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597316
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1208: acaatggatg agttataggt ttggttaaga cttgaaaagt cattgatcac ggcttttgtc cattattccg tgtttaaagt aatttacatt ctcttttaag atctctcagt ccaaagacaa 120 tgcaaacaat aataatttct ccacttgttt ctcatcgtct ctgtcttgct cgtgctgttc 180 cttgcaaccg tcttctcaac aaccaccacc gtgctcctcc ttcgatccgc ctctcaaacc 240 300 acceptica caccitication catches catches agree agree agreement agre tggcgactga agatgtgcaa gatcccagaa tcgctaagat tgcctcttcc attagagtca 360 420 tccccgactt ccctaaacca ggaatcatgt ttcaggacat aacgacgctt cttctcgaca 480 ctgaggcctt taaggatact attgctttgt ttgttgatag atacaaagat aaaggcatat ctgttgttgc aggtgttgaa gctagaggtt tcatttttgg ccctcctatt gcgttggcta 540 600 ttggtgccaa atttgttccc atgaggacgc ccaagaagct acctgggaag gttatttcgg aggagtattc gttggagtat ggaccagata cgattgagat gcacgtaggt gcagtagagc 660 720 ctggtgagcg tgctattatt attgatgncc tcattgccac gggtgggact ctcgctgctg caatccgact acttgaacga gtaggagtgn agattgttga atgtgcttgc gtaattgagt 780 taccagaget taagggaaag gagaaactag gagagaegte getatttgtt ettgtaaagt 840 cggctgctta acaagaaact ggaagtgaag gttattggat cgagtgttga tgctattttt 900 catgtatggt gagacatttt gcgtgggatt tgatccttgt tgnttcaact tttatcatan 960 ttggttcagn ctagaaaatg gcatttgaat gtcaggattc gattgcagtt tcctattgtt 1020 ccacttaaaa tccccg
- (2) INFORMATION FOR SEQ ID NO:1209:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 243 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..243

(D) OTHER INFORMATION: / Ceres Seq. ID 1597317 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1209: Met Gln Thr Ile Ile Ile Ser Pro Leu Val Ser His Arg Leu Cys Leu 10 Ala Arg Ala Val Pro Cys Asn Arg Leu Leu Asn Asn His His Arg Ala 25 Pro Pro Ser Ile Arg Leu Ser Asn His Arg Ser Thr Thr Ser Leu Arg 40 45 Leu Phe Ser Ser Ala Ala Ala Ser Arg Asp Ser Glu Met Ala Thr Glu 55 60 Asp Val Gln Asp Pro Arg Ile Ala Lys Ile Ala Ser Ser Ile Arg Val 75 Ile Pro Asp Phe Pro Lys Pro Gly Ile Met Phe Gln Asp Ile Thr Thr 90 85 Leu Leu Leu Asp Thr Glu Ala Phe Lys Asp Thr Ile Ala Leu Phe Val 105 Asp Arg Tyr Lys Asp Lys Gly Ile Ser Val Val Ala Gly Val Glu Ala 120 Arg Gly Phe Ile Phe Gly Pro Pro Ile Ala Leu Ala Ile Gly Ala Lys 135 140 Phe Val Pro Met Arg Thr Pro Lys Lys Leu Pro Gly Lys Val Ile Ser 150 155 Glu Glu Tyr Ser Leu Glu Tyr Gly Pro Asp Thr Ile Glu Met His Val 170 165 Gly Ala Val Glu Pro Gly Glu Arg Ala Ile Ile Ile Asp Xaa Leu Ile 185 190 180 Ala Thr Gly Gly Thr Leu Ala Ala Ile Arg Leu Leu Glu Arg Val 200 Gly Val Xaa Ile Val Glu Cys Ala Cys Val Ile Glu Leu Pro Glu Leu 210 215 220

- (2) INFORMATION FOR SEQ ID NO:1210:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 183 amino acids

230

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:

Ser Ala Ala

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..183
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597318
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1210:
- Met Ala Thr Glu Asp Val Gln Asp Pro Arg Ile Ala Lys Ile Ala Ser 1 5 10 15

Lys Gly Lys Glu Lys Leu Gly Glu Thr Ser Leu Phe Val Leu Val Lys

- Ser Ile Arg Val Ile Pro Asp Phe Pro Lys Pro Gly Ile Met Phe Gln
- Asp Ile Thr Thr Leu Leu Leu Asp Thr Glu Ala Phe Lys Asp Thr Ile 35 40 45
- Ala Leu Phe Val Asp Arg Tyr Lys Asp Lys Gly Ile Ser Val Val Ala
- Gly Val Glu Ala Arg Gly Phe Ile Phe Gly Pro Pro Ile Ala Leu Ala 65 70 75 80
- Ile Gly Ala Lys Phe Val Pro Met Arg Thr Pro Lys Lys Leu Pro Gly
 85 90 95
- Lys Val Ile Ser Glu Glu Tyr Ser Leu Glu Tyr Gly Pro Asp Thr Ile

105 100 Glu Met His Val Gly Ala Val Glu Pro Gly Glu Arg Ala Ile Ile Ile 125 120 Asp Xaa Leu Ile Ala Thr Gly Gly Thr Leu Ala Ala Ala Ile Arg Leu 140 135 Leu Glu Arg Val Gly Val Xaa Ile Val Glu Cys Ala Cys Val Ile Glu 150 155 Leu Pro Glu Leu Lys Gly Lys Glu Lys Leu Gly Glu Thr Ser Leu Phe 170 165 Val Leu Val Lys Ser Ala Ala 180 (2) INFORMATION FOR SEQ ID NO:1211: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 154 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..154 (D) OTHER INFORMATION: / Ceres Seq. ID 1597319 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1211: Met Phe Gln Asp Ile Thr Thr Leu Leu Leu Asp Thr Glu Ala Phe Lys 10 5 Asp Thr Ile Ala Leu Phe Val Asp Arg Tyr Lys Asp Lys Gly Ile Ser 2.5 Val Val Ala Gly Val Glu Ala Arg Gly Phe Ile Phe Gly Pro Pro Ile 45 40 Ala Leu Ala Ile Gly Ala Lys Phe Val Pro Met Arg Thr Pro Lys Lys 55 Leu Pro Gly Lys Val Ile Ser Glu Glu Tyr Ser Leu Glu Tyr Gly Pro 75 70 Asp Thr Ile Glu Met His Val Gly Ala Val Glu Pro Gly Glu Arg Ala 90 85 Ile Ile Ile Asp Xaa Leu Ile Ala Thr Gly Gly Thr Leu Ala Ala 105 Ile Arg Leu Leu Glu Arg Val Gly Val Xaa Ile Val Glu Cys Ala Cys 120 Val Ile Glu Leu Pro Glu Leu Lys Gly Lys Glu Lys Leu Gly Glu Thr 135 Ser Leu Phe Val Leu Val Lys Ser Ala Ala 150 (2) INFORMATION FOR SEQ ID NO:1212: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 1097 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: -(B) LOCATION: 1..1097 (D) OTHER INFORMATION: / Ceres Seq. ID 1597344 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1212: cacaatttga tacgcgtcag tttgaccaga ggctgaatga agttctcgat ggacaggatg

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1212:

cacaatttga tacgcgtcag tttgaccaga ggctgaatga agttctcgat ggacaggatg

agttcttcac ctcatatgat gaggtccatg agagctttga tgccatgggt ttgcaagaga

atcttcttag gggtatctat gcttacggtt tcgaaaagcc ttctgctatt cagcaaagag

gaattgtacc cttttgcaag ggtcttgatg tgatccagca ggcacagtct ggtactggaa

aaaccgccac tttctgctct ggtgtcttgc agcagcttga ctatgccctt ctccagtgcc

aggctctcgt tttggctccc accagagagc ttgctcagca gattgagaag gtcatgcgtg

360

1020

1080

cccttggtga (ctaccaaggt	gtcaaggttc	atgcctgtgt	tggtggaacc	agtgtccgtn	420
aggatcagcg (cattctccag	gctggtgttc	atgttgtcgt	tggaactcct	ggtcgtgttt	480
ttgacatgct t	tagaagacaa	tctctccgcc	ctgactgcat	caagatgttt	gtccttgatg	540
aagctgatga a	aatgctctcc	cgtggtttca	aggatcagct	catggagaca	tggaccaaaa	600
caccagagac a	atcatcatga	gagagttcag	gtcntggctc	gtctcgtgtt	ctaatcacaa	660
ccgatctctt (ggctcgtggt	atcgatgtgc	agcaagtctc	tctagtcata	aanctttgac	720
ctcccaactc a	agccagagaa	ttaccttcac	cgtatcggaa	gaagtggaag	gttcgggaga	780
aagggtgtgg d	cgattaactt	tgtgancnnt	tgangatcag	agaatgctgt	ttgatattca	840
gaaantctnc a	aatgtnggtt	gttgagnagc	ttcccncaaa	cgtggctgat	ttngctgtga	900
agaagaagaa 🤄	gaagaagaan	gaagaaaggt	tngcttgttn	cgcnnntttc	ttatttngat	960

- (2) INFORMATION FOR SEQ ID NO:1213:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 205 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

caatatttat ttcttct

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..205
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597345

ttttgacccg tctnnnnntc tctgaatcat cgatacatct ttttcattct ctagtattct

tcatcaactc tatcgaacat atatatgtct cttctgacca tatttctttn ttatcaaatg

- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1213:
- Gln Phe Asp Thr Arg Gln Phe Asp Gln Arg Leu Asn Glu Val Leu Asp
 1 5 10 15
- Gly Gln Asp Glu Phe Phe Thr Ser Tyr Asp Glu Val His Glu Ser Phe 20 25 30
- Asp Ala Met Gly Leu Gln Glu Asn Leu Leu Arg Gly Ile Tyr Ala Tyr 35 40 45
- Gly Phe Glu Lys Pro Ser Ala Ile Gln Gln Arg Gly Ile Val Pro Phe 50 60
- Cys Lys Gly Leu Asp Val Ile Gln Gln Ala Gln Ser Gly Thr Gly Lys 65 70 75 80
 Thr Ala Thr Phe Cys Ser Gly Val Leu Gln Gln Leu Asp Tyr Ala Leu
- 85 90 95
 Leu Gln Cys Gln Ala Leu Val Leu Ala Pro Thr Arg Glu Leu Ala Gln
- 100 105 110
- Gln Ile Glu Lys Val Met Arg Ala Leu Gly Asp Tyr Gln Gly Val Lys 115 120 125
- Val His Ala Cys Val Gly Gly Thr Ser Val Arg Xaa Asp Gln Arg Ile 130 135 140
- Leu Gln Ala Gly Val His Val Val Val Gly Thr Pro Gly Arg Val Phe 145 150 155 160
- Asp Met Leu Arg Arg Gln Ser Leu Arg Pro Asp Cys Ile Lys Met Phe 165 170 175
- Val Leu Asp Glu Ala Asp Glu Met Leu Ser Arg Gly Phe Lys Asp Gln 180 185 190
- Leu Met Glu Thr Trp Thr Lys Thr Pro Glu Thr Ser Ser 195 200 205
- (2) INFORMATION FOR SEQ ID NO:1214:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 171 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..171

(D) OTHER INFORMATION: / Ceres Seq. ID 1597346 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1214: Met Gly Leu Gln Glu Asn Leu Leu Arg Gly Ile Tyr Ala Tyr Gly Phe 10 Glu Lys Pro Ser Ala Ile Gln Gln Arg Gly Ile Val Pro Phe Cys Lys 25 Gly Leu Asp Val Ile Gln Gln Ala Gln Ser Gly Thr Gly Lys Thr Ala 40 Thr Phe Cys Ser Gly Val Leu Gln Gln Leu Asp Tyr Ala Leu Leu Gln 60 55 Cys Gln Ala Leu Val Leu Ala Pro Thr Arg Glu Leu Ala Gln Gln Ile 70 75 Glu Lys Val Met Arg Ala Leu Gly Asp Tyr Gln Gly Val Lys Val His 90 85 Ala Cys Val Gly Gly Thr Ser Val Arg Xaa Asp Gln Arg Ile Leu Gln 110 105 100 Ala Gly Val His Val Val Val Gly Thr Pro Gly Arg Val Phe Asp Met 125 120 Leu Arg Arg Gln Ser Leu Arg Pro Asp Cys Ile Lys Met Phe Val Leu 135 140 Asp Glu Ala Asp Glu Met Leu Ser Arg Gly Phe Lys Asp Gln Leu Met 150 155 Glu Thr Trp Thr Lys Thr Pro Glu Thr Ser Ser 165 (2) INFORMATION FOR SEQ ID NO:1215:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 123 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..123
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597347
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1215:

Met Pro Phe Ser Ser Ala Arg Leu Ser Phe Trp Leu Pro Pro Glu Ser 10 5 Leu Leu Ser Arg Leu Arg Arg Ser Cys Val Pro Leu Val Thr Thr Lys

30 25 20 Val Ser Arg Phe Met Pro Val Leu Val Glu Pro Val Ser Xaa Arg Ile

45 40 Ser Ala Phe Ser Arg Leu Val Phe Met Leu Ser Leu Glu Leu Leu Val 60 5.5

Val Phe Leu Thr Cys Leu Glu Asp Asn Leu Ser Ala Leu Thr Ala Ser 75 70 Arg Cys Leu Ser Leu Met Lys Leu Met Lys Cys Ser Pro Val Val Ser

90 85 Arg Ile Ser Ser Trp Arg His Gly Pro Lys His Gln Arg His His His

100 105 Glu Arg Val Gln Val Xaa Ala Arg Leu Val Phe

- 120 (2) INFORMATION FOR SEQ ID NO:1216:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 1156 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -

(B) LOCATION: 1..1156

(D) OTHER INFORMATION: / Ceres Seq. ID 1597360

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1216: tgtgcggctc gaagaaggaa cggtgaagtc tttaatagaa gaaaaaggag tgatcaaagg 60 agtgacatac aaaaataaag caggcgaaga aacaacagcc ttggcacctc tcactgtagt 120 atgcgacggt tgctactcaa accttcgtcg gtctctaaac gataacaatg ctgaggttct 180 240 gtcttaccaa gttgcttaca tctcgaagaa ttgtcggctt gatgatccca aaaaccttca 300 tttgataatg tccaaaccct cctacatcat gttgtaccaa ttaagcagca ctgatgttcg ttgtggtttt gaacttttct ccgccaattt tccttccatt gcaaaaggtg aaatggctac 360 ttttgcgaag aacactctag ctcctcaggt acctccaaaa cttcgcaaaa tattcttgaa 420 aggtctagat gaaggagcac acataaaagc ggtgccagca aagcgcatga cagcttcttt 480 540 aaccaagaaa aatggagtga ttgtgttggg agatgcattc aacatgcgtc atcnttcggt 600 cgcggctggt atgatggttt tattgtctga cattctcatt ctacgtcgtc ttctccagcc actaagcaac cttggtgatg caaacaaagt ctcagaagtc attaagtcct ttaatattat 660 ccgcaagcca atgtcagcga cggtgaacac attaggaaat gcattttctc aagttctaat 720 tgcatcaact gacgaagcaa aagaggcaat gagacaaggt tgctatgatt acctctctag 780 tggtggtttt tgcacgtcgg ggatgatggc tctactaggt ggcatgaatc ctcgtccgct 840 ctctctcata tatcatctat gtgctatcac tttacattca attggccatc ttctatctcc 900 960 gttggttccc catctcaagg ctgagggagt tagcgaaatg tttttccag caaacgcagc 1020 cgcatatcgc aaaagctata tggccgcatc tggtctttaa acattgatgc tctaaacagc 1080 aacatacgta agcttgtaaa ctcgttgaaa tgttattgta atgcttagct tacataatat 1140 tacttaatga gtttct

- (2) INFORMATION FOR SEQ ID NO:1217:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 352 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..352
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597361
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1217: Val Arg Leu Glu Glu Gly Thr Val Lys Ser Leu Ile Glu Glu Lys Gly
- 20 25 30

 Ala Leu Ala Pro Leu Thr Val Val Cys Asp Gly Cys Tyr Ser Asn Leu
- 35 40 45 Arg Arg Ser Leu Asn Asp Asn Asn Ala Glu Val Leu Ser Tyr Gln Val
- 50 55 60
 Ala Tyr Ile Ser Lys Asn Cys Arg Leu Asp Asp Pro Lys Asn Leu His
- 65 70 75 80
 Leu Ile Met Ser Lys Pro Ser Tyr Ile Met Leu Tyr Gln Leu Ser Ser
- 85 90 95

 Thr Asp Val Arg Cys Gly Phe Glu Leu Phe Ser Ala Asn Phe Pro Ser
- 100 105 110

 Ile Ala Lys Gly Glu Met Ala Thr Phe Ala Lys Asn Thr Leu Ala Pro
 115 120 125
- Gln Val Pro Pro Lys Leu Arg Lys Ile Phe Leu Lys Gly Leu Asp Glu
 130 135 140
- Gly Ala His Ile Lys Ala Val Pro Ala Lys Arg Met Thr Ala Ser Leu 145 150 155 160
- Thr Lys Lys Asn Gly Val Ile Val Leu Gly Asp Ala Phe Asn Met Arg
- His Xaa Ser Val Ala Ala Gly Met Met Val Leu Leu Ser Asp Ile Leu 180 185 190
- Ile Leu Arg Arg Leu Leu Gln Pro Leu Ser Asn Leu Gly Asp Ala Asn 195 200 205

Lys Val Ser Glu Val Ile Lys Ser Phe Asn Ile Ile Arg Lys Pro Met 220 215 Ser Ala Thr Val Asn Thr Leu Gly Asn Ala Phe Ser Gln Val Leu Ile 230 235 Ala Ser Thr Asp Glu Ala Lys Glu Ala Met Arg Gln Gly Cys Tyr Asp 250 245 Tyr Leu Ser Ser Gly Gly Phe Cys Thr Ser Gly Met Met Ala Leu Leu 260 265 Gly Gly Met Asn Pro Arg Pro Leu Ser Leu Ile Tyr His Leu Cys Ala 280 285 Ile Thr Leu His Ser Ile Gly His Leu Leu Ser Pro Phe Pro Ser Pro 290 295 300 Leu Ser Ile Trp His Ser Leu Arg Leu Phe Gly Leu Ala Leu Lys Met 310 315 Leu Val Pro His Leu Lys Ala Glu Gly Val Ser Glu Met Phe Pro 325 330 Ala Asn Ala Ala Tyr Arg Lys Ser Tyr Met Ala Ala Ser Gly Leu 340 345

- (2) INFORMATION FOR SEQ ID NO:1218:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 270 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..270
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597362
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1218: Met Ser Lys Pro Ser Tyr Ile Met Leu Tyr Gln Leu Ser Ser Thr Asp 10 5 Val Arg Cys Gly Phe Glu Leu Phe Ser Ala Asn Phe Pro Ser Ile Ala 25 Lys Gly Glu Met Ala Thr Phe Ala Lys Asn Thr Leu Ala Pro Gln Val 4 0 Pro Pro Lys Leu Arg Lys Ile Phe Leu Lys Gly Leu Asp Glu Gly Ala 55 His Ile Lys Ala Val Pro Ala Lys Arg Met Thr Ala Ser Leu Thr Lys 75 70 Lys Asn Gly Val Ile Val Leu Gly Asp Ala Phe Asn Met Arg His Xaa 90 85 Ser Val Ala Ala Gly Met Met Val Leu Leu Ser Asp Ile Leu Ile Leu 100 105 Arg Arg Leu Leu Gln Pro Leu Ser Asn Leu Gly Asp Ala Asn Lys Val 120 125 Ser Glu Val Ile Lys Ser Phe Asn Ile Ile Arg Lys Pro Met Ser Ala 135 140 Thr Val Asn Thr Leu Gly Asn Ala Phe Ser Gln Val Leu Ile Ala Ser 150 . 155 Thr Asp Glu Ala Lys Glu Ala Met Arg Gln Gly Cys Tyr Asp Tyr Leu 170 175 165 Ser Ser Gly Gly Phe Cys Thr Ser Gly Met Met Ala Leu Leu Gly Gly 185 190 Met Asn Pro Arg Pro Leu Ser Leu Ile Tyr His Leu Cys Ala Ile Thr 200 205 Leu His Ser Ile Gly His Leu Leu Ser Pro Phe Pro Ser Pro Leu Ser

Ile Trp His Ser Leu Arg Leu Phe Gly Leu Ala Leu Lys Met Leu Val

210 215 220

Attorney Docket No. 750-1237P Client Docket No. 46.003 230 235 225 Pro His Leu Lys Ala Glu Gly Val Ser Glu Met Phe Pro Ala Asn 245 250 255 Ala Ala Ala Tyr Arg Lys Ser Tyr Met Ala Ala Ser Gly Leu 260 265 (2) INFORMATION FOR SEQ ID NO:1219: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 263 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..263 (D) OTHER INFORMATION: / Ceres Seq. ID 1597363 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1219: Met Leu Tyr Gln Leu Ser Ser Thr Asp Val Arg Cys Gly Phe Glu Leu 5 10 Phe Ser Ala Asn Phe Pro Ser Ile Ala Lys Gly Glu Met Ala Thr Phe 2.5 Ala Lys Asn Thr Leu Ala Pro Gln Val Pro Pro Lys Leu Arg Lys Ile 4 0 Phe Leu Lys Gly Leu Asp Glu Gly Ala His Ile Lys Ala Val Pro Ala 55 Lys Arg Met Thr Ala Ser Leu Thr Lys Lys Asn Gly Val Ile Val Leu 70 Gly Asp Ala Phe Asn Met Arg His Xaa Ser Val Ala Ala Gly Met Met 90 85 Val Leu Leu Ser Asp Ile Leu Ile Leu Arg Arg Leu Leu Gln Pro Leu 105 110

Ser Asn Leu Gly Asp Ala Asn Lys Val Ser Glu Val Ile Lys Ser Phe 115 120 125 Asn Ile Ile Arg Lys Pro Met Ser Ala Thr Val Asn Thr Leu Gly Asn 130 135 140 Ala Phe Ser Gln Val Leu Ile Ala Ser Thr Asp Glu Ala Lys Glu Ala 155 150 Met Arg Gln Gly Cys Tyr Asp Tyr Leu Ser Ser Gly Gly Phe Cys Thr 165 170 Ser Gly Met Met Ala Leu Leu Gly Gly Met Asn Pro Arg Pro Leu Ser 180 185 190 Leu Ile Tyr His Leu Cys Ala Ile Thr Leu His Ser Ile Gly His Leu 195 200 205 Leu Ser Pro Phe Pro Ser Pro Leu Ser Ile Trp His Ser Leu Arg Leu 210 215 220 Phe Gly Leu Ala Leu Lys Met Leu Val Pro His Leu Lys Ala Glu Gly 230 235 Val Ser Glu Met Phe Phe Pro Ala Asn Ala Ala Ala Tyr Arg Lys Ser 250 245 Tyr Met Ala Ala Ser Gly Leu

(2) INFORMATION FOR SEQ ID NO:1220:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 504 base pairs
 - (B) TYPE: nucleic acid

 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..504

(D) OTHER INFORMATION: / Ceres Seq. ID 1597376

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1220: 60 aatcacggcg gcggcggcgg aagagtagca gcagcaggcg gcgccgagta gcgtctcccc acctcgagct tgccaccatg gctagaggat tgaagaagca tttgaagagg ctcaatgcgc 120 ccaagcattg gatgctggac aagcttggcg gasttttgct cccaaaccat cttctggacc 180 tcacaagtct agggagtgcc tgccactgat cctcatcatc aggaacaggc tcaagtatgc 240 tettacatac egtgagteat ttecateetg atgeaacgee atgtacttgt tgatggeaag 300 gtcaggacag acaagaccta ccctgctggg ttcatggatg tcatttccat ccccaagacc 360 aacgagaact acaggctgct gtatgacact aagggccgct tecgeettea eecaatcagg 420 gatgaggatg ctaagttcaa gctttgcaag gttaggtctg ttcagtttgg gcagaagggc 480 atcccqtacc tgaacacgta tgac

- (2) INFORMATION FOR SEQ ID NO:1221:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 68 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..68
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597377

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1221:

Ser Arg Arg Arg Arg Lys Ser Ser Ser Ser Arg Arg Arg Val $1 \ 5 \ 10 \ 15$ Ala Ser Pro His Leu Glu Leu Ala Thr Met Ala Arg Gly Leu Lys Lys

20 25 30

His Leu Lys Arg Leu Asn Ala Pro Lys His Trp Met Leu Asp Lys Leu 35 40 45

Gly Gly Xaa Leu Leu Pro Asn His Leu Leu Asp Leu Thr Ser Leu Gly 50 55 60

Ser Ala Cys His

65

- (2) INFORMATION FOR SEQ ID NO:1222:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 78 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..78
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597378
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1222:

Met Gln Arg His Val Leu Val Asp Gly Lys Val Arg Thr Asp Lys Thr

1 10 15

Tyr Pro Ala Gly Phe Met Asp Val Ile Ser Ile Pro Lys Thr Asn Glu 20 25 30

Asn Tyr Arg Leu Leu Tyr Asp Thr Lys Gly Arg Phe Arg Leu His Pro 35 40 45

Ile Arg Asp Glu Asp Ala Lys Phe Lys Leu Cys Lys Val Arg Ser Val

Gln Phe Gly Gln Lys Gly Ile Pro Tyr Leu Asn Thr Tyr Asp
65 70 75

- (2) INFORMATION FOR SEQ ID NO:1223:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 70 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: peptide (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..70
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597379

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1223: Met Tyr Leu Leu Met Ala Arg Ser Gly Gln Thr Arg Pro Thr Leu Leu 10 5 Gly Ser Trp Met Ser Phe Pro Ser Pro Arg Pro Thr Arg Thr Thr Gly 30 25 20

Cys Cys Met Thr Leu Arg Ala Ala Ser Ala Phe Thr Gln Ser Gly Met 40 45

Arg Met Leu Ser Ser Ser Phe Ala Arg Leu Gly Leu Phe Ser Leu Gly 60 55

Arg Arg Ala Ser Arg Thr 70

- (2) INFORMATION FOR SEQ ID NO:1224:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 466 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..466
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597392
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1224: aatageetee geeteteett eeagtgteeg eegtegtgeg etegetaeee eteteeeteg 60 120 aggectttge eggegaagag egeceagteg eccaceagga tgaagtttgt tgetgeetae ctgcttgctg tcctcgctgg gaactccagc ccctctgccg aggacttgac agccattctg 180 gagtcagttg gctgtgaagt tgacaatgaa aagatggaac tccttctgtc ccaactgagc 240 ggtaaggaca ttaccgagct cattgctgct ggcagggaga agtttgcttc agtcccatgt 300 ggcggtgggc ggtgtggctg ttgtggcagc tgcccctgct gctggcggcg ctcctgcagc 360 tgaggcgaag aaagaagaga aggtggagga gaaggaagaa agtgatgacg acatgggctt 420 cagoctotto gactaagoot gtgcatagto aagagtattg tttttg
- (2) INFORMATION FOR SEQ ID NO:1225:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 120 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..120
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597393
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1225:

Asn Ser Leu Arg Leu Ser Phe Gln Cys Pro Pro Ser Cys Ala Arg Tyr 10 5 Pro Ser Pro Ser Arg Pro Leu Pro Ala Lys Ser Ala Gln Ser Pro Thr

25 Arg Met Lys Phe Val Ala Ala Tyr Leu Leu Ala Val Leu Ala Gly Asn

40

Ser Ser Pro Ser Ala Glu Asp Leu Thr Ala Ile Leu Glu Ser Val Gly 55

Cys Glu Val Asp Asn Glu Lys Met Glu Leu Leu Ser Gln Leu Ser 75 70 65

Gly Lys Asp Ile Thr Glu Leu Ile Ala Ala Gly Arg Glu Lys Phe Ala 90 Ser Val Pro Cys Gly Gly Gly Arg Cys Gly Cys Cys Gly Ser Cys Pro

100 105 110

Cys Cys Trp Arg Arg Ser Cys Ser 115 120

- (2) INFORMATION FOR SEQ ID NO:1226:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 87 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..87
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597394
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1226: Met Lys Phe Val Ala Ala Tyr Leu Leu Ala Val Leu Ala Gly Asn Ser

1 5 10 15
Ser Pro Ser Ala Glu Asp Leu Thr Ala Ile Leu Glu Ser Val Gly Cys
20 25 30

Glu Val Asp Asn Glu Lys Met Glu Leu Leu Ser Gln Leu Ser Gly

Lys Asp Ile Thr Glu Leu Ile Ala Ala Gly Arg Glu Lys Phe Ala Ser
50 55 60

Val Pro Cys Gly Gly Gly Arg Cys Gly Cys Cys Gly Ser Cys Pro Cys
65 70 75 80

Cys Trp Arg Arg Ser Cys Ser

85

- (2) INFORMATION FOR SEQ ID NO:1227:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 502 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..502
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597396
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1227:

aatageetee geeteteett eeagtgteeg eegtegtgeg etegetaeee eteteeeteg 60 aggeetttge egegaagage geceagtege ceaceaggat gaagtttgtt getgeetate 120 tgcttgctgt cctcgctggg aactccagcc cctctgccga ggacttgaca tccattctgg 180 agtcagttgg ctgtgaagtt gacaatgaaa agatggaact ccttctgtcc cagctgagcg 240 gtaaggacat taccgagctc attgccgccg gcagggagaa gtttgcttca gtcccatgtg 300 geggtggegg tgtggctgtc geggcastgc ccctgctgct ggcggcggcg gegetcctgc 360 agctgaggcg aagaaagaag agaaggtgga ggagaaggaa gaaagtgatg acgacatggg 420 cttcagctct tcgactaagt ctgtgcaata gccaagggta ttgtttttga gtcgaggaag 480

- cagaggggaa gaaaaatagt ag (2) INFORMATION FOR SEQ ID NO:1228:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 64 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..64
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597397
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1228:
- Ile Ala Ser Ala Ser Pro Ser Ser Val Arg Arg Arg Ala Leu Ala Thr

1 5 10 15
Pro Leu Pro Arg Gly Leu Cys Arg Glu Glu Arg Pro Val Ala His Gln
20 25 30

Asp Glu Val Cys Cys Cys Leu Ser Ala Cys Cys Pro Arg Trp Glu Leu 35 40 45

Gln Pro Leu Cys Arg Gly Leu Asp Ile His Ser Gly Val Ser Trp Leu 50 55 60

(2) INFORMATION FOR SEQ ID NO:1229:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 123 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..123
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597398
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1229:

Met Lys Phe Val Ala Ala Tyr Leu Leu Ala Val Leu Ala Gly Asn Ser 1 10 15

Ser Pro Ser Ala Glu Asp Leu Thr Ser Ile Leu Glu Ser Val Gly Cys 20 25 30

Glu Val Asp Asn Glu Lys Met Glu Leu Leu Leu Ser Gln Leu Ser Gly 35 40 45

Lys Asp Ile Thr Glu Leu Ile Ala Ala Gly Arg Glu Lys Phe Ala Ser 50 55 60

Val Pro Cys Gly Gly Gly Val Ala Val Ala Ala Xaa Pro Leu Leu 65 70 75 80

Leu Ala Ala Ala Leu Leu Gln Leu Arg Arg Arg Lys Lys Arg Arg 85 90 95

Trp Arg Arg Lys Lys Val Met Thr Trp Ala Ser Ala Leu Arg
100 105 110

Leu Ser Leu Cys Asn Ser Gln Gly Tyr Cys Phe 115 120

- (2) INFORMATION FOR SEQ ID NO:1230:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 85 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..85
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597399

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1230:

Met Glu Leu Leu Ser Gln Leu Ser Gly Lys Asp Ile Thr Glu Leu
1 5 10 15

Ile Ala Ala Gly Arg Glu Lys Phe Ala Ser Val Pro Cys Gly Gly Gly 20 25 30

Gly Val Ala Val Ala Ala Xaa Pro Leu Leu Leu Ala Ala Ala Leu 35 40 45

Leu Gln Leu Arg Arg Arg Lys Lys Arg Arg Trp Arg Arg Arg Lys Lys 50 55 60

Val Met Thr Trp Ala Ser Ala Leu Arg Leu Ser Leu Cys Asn Ser 65 70 75 80,

Gln Gly Tyr Cys Phe

- (2) INFORMATION FOR SEQ ID NO:1231:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 509 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..509
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597412
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1231:

60 cacccaagte caaqcaccdn teccaaaaaa atececcatt ttettegagg egaggagage gtgaaagaca gggcaccgnc agcgacgatg tcggggcgcg gcaagggcgg caaaggtctg 120 ggcaagggcg gasaaagcgc caccgtaagt ctccgtgaca acatccaggr gcatcaccaa 180 gcccgcgatc cgtagctggc tcggaggggc ggcgtgaagc sattttcggg gcttatctac 240 gaggagaccc gcggcgtcct caagatcttc ctcgagaacg tcatccgcga cgccgtcacc 300 tacaccgage acgeacgecg caagactgtt accgccatgg acgtcgtcta cgcgctcaag 360 cscagggccg aaccetetae ggetteggag getaggetgg atgeetette teettgetgt 420 cgtggatgcc tgtttgttgt tagggttact tcgtttagtt ccttgtctgt gacgtgttgg 480 actattgtta gtagtattct tgaagttgt

- (2) INFORMATION FOR SEQ ID NO:1232:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 64 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..64
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597413
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1232:

Gly Glu Glu Ser Val Lys Asp Arg Ala Pro Xaa Ala Thr Met Ser Gly 20 25 30

Arg Gly Lys Gly Gly Lys Gly Leu Gly Lys Gly Gly Xaa Ser Ala Thr

Val Ser Leu Arg Asp Asn Ile Gln Xaa His His Gln Ala Arg Asp Pro 50 55 60

- (2) INFORMATION FOR SEQ ID NO:1233:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 51 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..51
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597414
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1233:

Pro Lys Ser Lys His Xaa Ser Gln Lys Asn Pro Pro Phe Ser Ser Arg

Arg Gly Glu Arg Glu Arg Gln Gly Thr Xaa Ser Asp Asp Val Gly Ala

Arg Gln Gly Arg Gln Arg Ser Gly Gln Gly Arg Xaa Lys Arg His Arg

Lys Ser Pro 50

- (2) INFORMATION FOR SEQ ID NO:1234:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 57 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..57
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597415
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1234:

Met Asp Val Val Tyr Ala Leu Lys Xaa Arg Ala Glu Pro Ser Thr Ala 1 10 15

Ser Glu Ala Arg Leu Asp Ala Ser Ser Pro Cys Cys Arg Gly Cys Leu 20 25 30

Phe Val Val Arg Val Thr Ser Phe Ser Ser Leu Ser Val Thr Cys Trp 35 40 45

Thr Ile Val Ser Ser Ile Leu Glu Val

- (2) INFORMATION FOR SEQ ID NO:1235:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 509 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..509
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597416
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1235:

tngctctcaa atatctctca ccacctctss sctcagcttn tcsscacctt tggtcaggca 60 ggctcaggca aagggggcgg msaacaagga cgagsgatgg sgaccaaggt catctgcttc 120 cttgtgctcg catccctgct cctcgccgtc gcctttcccg tgtctgctct gcgncagcrg 180 ttcgaagaag ggcggcggcg gcggtggtgg cggaggcagt gttagcggaa sgcaggcgcc 240 300 aacctgaatc cctgggagtg ctcgcccaag tgcgggtcgc ggtgctccaa gacgcagtac aggaaggcct gcctcacctt atgcaacaag tgctgcgcca agtgcctctg cgtgccaccg 360 gggttctacg gcaacaaggg cgcctgccct gctacaacaa ctggaaaacc aaggaaggag 420 ggcccaagtg cccctagaat aagatccacc gcagctcccg tccgccattg tcscmccttc 480 tccqaatctq qqaacgtgtt gttcatctt

- (2) INFORMATION FOR SEQ ID NO:1236:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 74 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..74
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597417
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1236:

Xaa Leu Ser Asn Ile Ser His His Leu Xaa Xaa Gln Leu Xaa Xaa Thr 1 5 10 15

Phe Gly Gln Ala Gly Ser Gly Lys Gly Gly Xaa Xaa Gln Gly Arg Xaa 20 25 30

Met Xaa Thr Lys Val Ile Cys Phe Leu Val Leu Ala Ser Leu Leu Leu 35 40 45

Ala Val Ala Phe Pro Val Ser Ala Leu Xaa Gln Xaa Phe Glu Glu Gly 50 55 60

Arg Arg Arg Trp Trp Arg Arg Gln Cys 70

- (2) INFORMATION FOR SEQ ID NO:1237:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 81 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..81
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597418
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1237:

Xaa Ser Gln Ile Ser Leu Thr Thr Ser Xaa Leu Ser Xaa Xaa Pro

Leu Val Arg Gln Ala Gln Ala Lys Gly Ala Xaa Asn Lys Asp Glu Xaa 20 25 30

Trp Xaa Pro Arg Ser Ser Ala Ser Leu Cys Ser His Pro Cys Ser Ser 35 40 45

Pro Ser Pro Phe Pro Cys Leu Leu Cys Xaa Ser Xaa Ser Lys Lys Gly

Gly Gly Gly Gly Gly Gly Gly Ser Val Ser Gly Xaa Gln Ala Pro 65 70 75 80

Thr

- (2) INFORMATION FOR SEQ ID NO:1238:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 144 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..144
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597419
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1238:

Ala Leu Lys Tyr Leu Ser Pro Pro Xaa Xaa Ser Ala Xaa Xaa His Leu 1 5 10 15

Trp Ser Gly Arg Leu Arg Gln Arg Gly Arg Xaa Thr Arg Thr Xaa Asp 20 25 30

Xaa Asp Gln Gly His Leu Leu Pro Cys Ala Arg Ile Pro Ala Pro Arg

Arg Arg Leu Ser Arg Val Cys Ser Ala Xaa Xaa Val Arg Arg Arg Ala 50 55 60

Ala Ala Val Val Ala Glu Ala Val Leu Ala Glu Xaa Arg Arg Gln
65 70 75 80

Pro Glu Ser Leu Gly Val Leu Ala Gln Val Arg Val Ala Val Leu Gln 85 90 95

Asp Ala Val Gln Glu Gly Leu Pro His Leu Met Gln Gln Val Leu Arg 100 105 110

Gln Val Pro Leu Arg Ala Thr Gly Val Leu Arg Gln Gln Gly Arg Leu 115 120 125

Pro Cys Tyr Asn Asn Trp Lys Thr Lys Glu Gly Gly Pro Lys Cys Pro 130 135 140

(2) INFORMATION FOR SEQ ID NO:1239:

80

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 492 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..492
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597426
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1239: aattttagga cctaattata agastctctc ttggagatgc tctaaggcac cgactcgccg 120 gggacagaga gagagaccga cgcgccggca gatcccaagc tcaccctaga gggggaagag ggcgaccgaa gcggcgatgg gtttcatcat ggacttcgcg gagaatctga tcctccgtct 180 240 gatggaggac ccggacaagc gcgaccaggt tcggcgggag catgtctaca agatgaagga 300 gcggtgcgag cgcactaagg cggcgtggag cctccctctg cgcccctacg gcttctggac 360 cttcgaccgc ttcaactcgc agctctcctg ggatccccag atcagccagg ccgccggcgt negggacece tacgacgace teategeacg ceaetetgge tegeogecgt ettectgaac 420 accogttccg atctttgccc agaaggtcta cttgggcatc aataagaaac tctttcccct 480
- (2) INFORMATION FOR SEQ ID NO:1240:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 93 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

caaactgatt gt

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..93
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597427
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1240:

Met Gly Phe Ile Met Asp Phe Ala Glu Asn Leu Ile Leu Arg Leu Met 1 5 10 15 Glu Asp Pro Asp Lys Arg Asp Gln Val Arg Arg Glu His Val Tyr Lys

20 25 30
Met Lys Glu Arg Cys Glu Arg Thr Lys Ala Ala Trp Ser Leu Pro Leu

Met Lys Giu Arg Cys Giu Arg ini Lys Aia Aia iip Sei hed iio hed 35 40 45

Arg Pro Tyr Gly Phe Trp Thr Phe Asp Arg Phe Asn Ser Gln Leu Ser 50 55 60

Trp Asp Pro Gln Ile Ser Gln Ala Ala Gly Xaa Arg Asp Pro Tyr Asp

Asp Leu Ile Ala Arg His Ser Gly Ser Pro Pro Ser Ser 85 90

- (2) INFORMATION FOR SEQ ID NO:1241:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 89 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..89
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597428
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1241:
- Met Asp Phe Ala Glu Asn Leu Ile Leu Arg Leu Met Glu Asp Pro Asp 1 5 10 15
- Lys Arg Asp Gln Val Arg Arg Glu His Val Tyr Lys Met Lys Glu Arg 20 25 30
- Cys Glu Arg Thr Lys Ala Ala Trp Ser Leu Pro Leu Arg Pro Tyr Gly

40 45

25 40 45
Phe Trp Thr Phe Asp Arg Phe Asn Ser Gln Leu Ser Trp Asp Pro Gln 50 55 60

Ile Ser Gln Ala Ala Gly Xaa Arg Asp Pro Tyr Asp Asp Leu Ile Ala 65 70 75 80

Arg His Ser Gly Ser Pro Pro Ser Ser 85

- (2) INFORMATION FOR SEQ ID NO:1242:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 78 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..78
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597429
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1242:
 Met Glu Asp Pro Asp Lys Arg Asp Gln Val Arg Arg Glu His Val Tyr

1 5 10 15 Lys Met Lys Glu Arg Cys Glu Arg Thr Lys Ala Ala Trp Ser Leu Pro

Lys Met Lys Glu Arg Cys Glu Arg Thr Lys Ala Ala Trp Ser Leu Pro 20 25 30

Leu Arg Pro Tyr Gly Phe Trp Thr Phe Asp Arg Phe Asn Ser Gln Leu 35 40 45

Ser Trp Asp Pro Gln Ile Ser Gln Ala Ala Gly Xaa Arg Asp Pro Tyr 50 60

Asp Asp Leu Ile Ala Arg His Ser Gly Ser Pro Pro Ser Ser 65 70 75

- (2) INFORMATION FOR SEQ ID NO:1243:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 868 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..868
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597430
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1243:

60 agaagettat egettateat eegegeetet acatttette tetetetget eeteegeete 120 cqcctccqsc tqqwtccsat ccaagacagg gatcgccatg tcggactcgg aggagcacca 180 cttcgaatcg aaggccgacg ctggggcgtc caagacctac ccgcagcagg ctggcaccgt 240 ccqtaagaac ggcttcatcg tcatcaagaa ccgtccctgc aaggttgtgg aggtttctac 300 ctccaagact ggtaagcatg gccacgccaa atgccacttt gtcgccatag acatattcaa 360 tgggaaaaag cttgaagata ttgttccttc atcacacaac tgtgacattc cgcatgtgaa 420 ccgtactgag tatcagctga ttgatatttc tgaggatgga tttgtgagcc ttcttacttc agatggcaac actaaggatg atcttagact cccaactgat gagactcttg tggcccagat 480 taaggaaggg ttcgaaagcg ggaaggatct tgttgtgact gtccagtctg cgatggggga 540 ggagcagatc tgcgcgctga aggatgttgg ccccaagtaa cttccttggc tttggagacg 600 gaaccctact ctatctcaac ttaaatcgga aaagaagtgt atcaccaaaa ggattgctac 660 cgagacattc acctggcttg agctbgcttt tgctatagcg tgcaaggggt tgctgggtgt 720 cgtacctacc tgacttatca ttatctgtgg ttacattctg gaccctgtat tttctatgct 780 840 ttcatgtctc tctactttct agtattaatt atcaataagt tggtttggaa tggttgaatt

(2) INFORMATION FOR SEQ ID NO:1244:

gcatccaatg catgaagttt ctttggct

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 192 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: peptide (B) LOCATION: 1..192

(D) OTHER INFORMATION: / Ceres Seq. ID 1597431

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1244: Glu Ala Tyr Arg Leu Ser Ser Ala Pro Leu His Phe Phe Ser Leu Cys 10 Ser Ser Ala Ser Ala Ser Xaa Trp Xaa Xaa Ser Lys Thr Gly Ile Ala 25 Met Ser Asp Ser Glu Glu His His Phe Glu Ser Lys Ala Asp Ala Gly 40 Ala Ser Lys Thr Tyr Pro Gln Gln Ala Gly Thr Val Arg Lys Asn Gly 55 Phe Ile Val Ile Lys Asn Arg Pro Cys Lys Val Val Glu Val Ser Thr 75 70 Ser Lys Thr Gly Lys His Gly His Ala Lys Cys His Phe Val Ala Ile 90 85 Asp Ile Phe Asn Gly Lys Lys Leu Glu Asp Ile Val Pro Ser Ser His 105 100 Asn Cys Asp Ile Pro His Val Asn Arg Thr Glu Tyr Gln Leu Ile Asp 120 125 Ile Ser Glu Asp Gly Phe Val Ser Leu Leu Thr Ser Asp Gly Asn Thr 135 140 Lys Asp Asp Leu Arg Leu Pro Thr Asp Glu Thr Leu Val Ala Gln Ile 155 150 Lys Glu Gly Phe Glu Ser Gly Lys Asp Leu Val Val Thr Val Gln Ser

170 Ala Met Gly Glu Glu Gln Ile Cys Ala Leu Lys Asp Val Gly Pro Lys

185

(2) INFORMATION FOR SEQ ID NO:1245:

180

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 160 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..160
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597432
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1245:

Met Ser Asp Ser Glu Glu His His Phe Glu Ser Lys Ala Asp Ala Gly 10

Ala Ser Lys Thr Tyr Pro Gln Gln Ala Gly Thr Val Arg Lys Asn Gly 25

Phe Ile Val Ile Lys Asn Arg Pro Cys Lys Val Val Glu Val Ser Thr

Ser Lys Thr Gly Lys His Gly His Ala Lys Cys His Phe Val Ala Ile 60

Asp Ile Phe Asn Gly Lys Lys Leu Glu Asp Ile Val Pro Ser Ser His 75 70

Asn Cys Asp Ile Pro His Val Asn Arg Thr Glu Tyr Gln Leu Ile Asp 90

Ile Ser Glu Asp Gly Phe Val Ser Leu Leu Thr Ser Asp Gly Asn Thr 105 110

Lys Asp Asp Leu Arg Leu Pro Thr Asp Glu Thr Leu Val Ala Gln Ile 120 125 115

60

120 180

Lys Glu Gly Phe Glu Ser Gly Lys Asp Leu Val Val Thr Val Gln Ser 140 135

Ala Met Gly Glu Glu Gln Ile Cys Ala Leu Lys Asp Val Gly Pro Lys 155 145 150

- (2) INFORMATION FOR SEQ ID NO:1246:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 239 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..239
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597437
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1246: acagcatete aagtetteae acteetegee ateacataaa accagtgeaa geagaageag cgcaatggcg agcagcacca tggccctctc ctccacagcc ttcgccggca akgcagtgaa cgtgccgtcg tctctcwtcg gcgangcccg cgtgacgatg cgcaagangg cggcgaakgc aaagccggcg gcgagctccg gcagcccgtg gtacggcccc gamcgcgtgc tctacctgg (2) INFORMATION FOR SEQ ID NO:1247:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 79 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..79
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597438
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1247:
- Gln His Leu Lys Ser Ser His Ser Ser Pro Ser His Lys Thr Ser Ala 15 5 10
- Ser Arg Ser Ser Ala Met Ala Ser Ser Thr Met Ala Leu Ser Ser Thr 25 30 20
- Ala Phe Ala Gly Xaa Ala Val Asn Val Pro Ser Ser Leu Xaa Gly Xaa 40 4.5
- Ala Arg Val Thr Met Arg Lys Xaa Ala Ala Xaa Ala Lys Pro Ala Ala 55 60
- Ser Ser Gly Ser Pro Trp Tyr Gly Pro Xaa Arg Val Leu Tyr Leu 70 75
- (2) INFORMATION FOR SEQ ID NO:1248:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 58 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..58
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597439
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1248:
- Met Ala Ser Ser Thr Met Ala Leu Ser Ser Thr Ala Phe Ala Gly Xaa 10 15 5 1
- Ala Val Asn Val Pro Ser Ser Leu Xaa Gly Xaa Ala Arg Val Thr Met 25
- Arg Lys Xaa Ala Ala Xaa Ala Lys Pro Ala Ala Ser Ser Gly Ser Pro

45 35 40 Trp Tyr Gly Pro Xaa Arg Val Leu Tyr Leu 55 50 (2) INFORMATION FOR SEQ ID NO:1249: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 53 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..53 (D) OTHER INFORMATION: / Ceres Seq. ID 1597440 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1249: Met Ala Leu Ser Ser Thr Ala Phe Ala Gly Xaa Ala Val Asn Val Pro 10 Ser Ser Leu Xaa Gly Xaa Ala Arg Val Thr Met Arg Lys Xaa Ala Ala 25 Xaa Ala Lys Pro Ala Ala Ser Ser Gly Ser Pro Trp Tyr Gly Pro Xaa 40 Arg Val Leu Tyr Leu 50 (2) INFORMATION FOR SEQ ID NO:1250: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 130 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: -(B) LOCATION: 1..130 (D) OTHER INFORMATION: / Ceres Seq. ID 1597452 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1250: tctcccaaag ccatcttcct tccgcggact agttttttct gaagattcgg cgatggcggc 60 gacgatccaa tccgtgaagg cgcgccagat ctttgacagc cgcggcaacc ctacagtcga 120 ggtcgacgtg (2) INFORMATION FOR SEQ ID NO:1251: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 43 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..43 (D) OTHER INFORMATION: / Ceres Seq. ID 1597453 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1251: Leu Pro Lys Pro Ser Ser Phe Arg Gly Leu Val Phe Ser Glu Asp Ser 15 5 10 Ala Met Ala Ala Thr Ile Gln Ser Val Lys Ala Arg Gln Ile Phe Asp 2.5 Ser Arg Gly Asn Pro Thr Val Glu Val Asp Val 40 35 (2) INFORMATION FOR SEQ ID NO:1252: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 446 base pairs

(B) TYPE: nucleic acid(C) STRANDEDNESS: single

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: DNA (genomic)

(ix) FEATURE:

- (A) NAME/KEY: -
- (B) LOCATION: 1..446
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597456
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1252: aagctcctcg tcaccagtca gaaaacagtg ggatcgagtt gtttcactgc acgagcacat 60 120 ceteeggega ceaeeggeet eeeteteegt eetetagega eegaeeaaeg egtegagega agatgtcgtg gcagacgtac gtggacgagc aaccctggat gtgcgagatc gagggccacc 180 acctcacgtc ggcggccatc gtcggccacg rgcgccacct gggctcagag caccgcattc 240 cccgagttca agcccgagga gatggctgcc atcatgaagg atttcgacga gccggggcac 300 tegeceegae eggeetgata etgggaggea ceaagtacat ggteateeaa gtegaacetg 360 gagctgtcat ccgtggcaag aagggatccg ggggcatcac tgtgaagaaa acagggcagt 420 cactcatcat tggcatctac gacgag
- (2) INFORMATION FOR SEQ ID NO:1253:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 105 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..105
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597457
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1253:
- Lys Leu Leu Val Thr Ser Gln Lys Thr Val Gly Ser Ser Cys Phe Thr 1 5 10 15
- Ala Arg Ala His Pro Pro Ala Thr Thr Gly Leu Pro Leu Arg Pro Leu 20 25 30
- Ala Thr Asp Gln Arg Val Glu Arg Arg Cys Arg Gly Arg Arg Thr Trp 35 40 45
- Thr Ser Asn Pro Gly Cys Ala Arg Ser Arg Ala Thr Thr Ser Arg Arg 50 55 60
- Arg Pro Ser Ser Ala Thr Xaa Ala Thr Trp Ala Gln Ser Thr Ala Phe 65 70 75 80
- Pro Glu Phe Lys Pro Glu Glu Met Ala Ala Ile Met Lys Asp Phe Asp 85 90 95
- Glu Pro Gly His Ser Pro Arg Pro Ala 100 105
- (2) INFORMATION FOR SEQ ID NO:1254:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 148 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..148
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597458
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1254:
- Ala Pro Arg His Gln Ser Glu Asn Ser Gly Ile Glu Leu Phe His Cys
 1 10 15
- Thr Ser Thr Ser Ser Gly Asp His Arg Pro Pro Ser Pro Ser Ser Ser 20 25 30
- Asp Arg Pro Thr Arg Arg Ala Lys Met Ser Trp Gln Thr Tyr Val Asp 35 40 45
- Glu Gln Pro Trp Met Cys Glu Ile Glu Gly His His Leu Thr Ser Ala
 50 55 60

Ala Ile Val Gly His Xaa Arg His Leu Gly Ser Glu His Arg Ile Pro
65 70 75 80

Arg Val Gln Ala Arg Gly Asp Gly Cys His His Glu Gly Phe Arg Arg
85 90 95

Ala Gly Ala Leu Ala Pro Thr Gly Leu Ile Leu Gly Gly Thr Lys Tyr
100 105 110

Met Val Ile Gln Val Glu Pro Gly Ala Val Ile Arg Gly Lys Lys Gly 115 120 125

Ser Gly Gly Ile Thr Val Lys Lys Thr Gly Gln Ser Leu Ile Ile Gly

Ile Tyr Asp Glu

145

- (2) INFORMATION FOR SEQ ID NO:1255:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 108 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..108
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597459
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1255:

Met Ser Trp Gln Thr Tyr Val Asp Glu Gln Pro Trp Met Cys Glu Ile 1 10 15

Glu Gly His His Leu Thr Ser Ala Ala Ile Val Gly His Xaa Arg His
20 25 30

Leu Gly Ser Glu His Arg Ile Pro Arg Val Gln Ala Arg Gly Asp Gly 35 40 45

Cys His His Glu Gly Phe Arg Arg Ala Gly Ala Leu Ala Pro Thr Gly 50 55 60

Leu Ile Leu Gly Gly Thr Lys Tyr Met Val Ile Gln Val Glu Pro Gly 65 70 75 80
Ala Val Ile Arg Gly Lys Lys Gly Ser Gly Gly Ile Thr Val Lys Lys

105

85 90
Thr Gly Gln Ser Leu Ile Ile Gly Ile Tyr Asp Glu

(2) INFORMATION FOR SEQ ID NO:1256:

100

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 472 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..472
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597476
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1256:

atetcageag tgacgaacte gaagettgek ttettgtagt agaggtegge ageaageggt 60 ggtagagega gaggtegga agatgaagag cageactetg ttggegatee tagttetea 120 ggeecttetg gtetctgegg eegtggeaaa ggacetgeag ggeegaegae gaagaageag 180 tgetggttge ggegagtgea ecasetggte ggeggtgtgg acetgegaeg aceteeteae 240 gaagtgegee geeaeetgea agaactgegt eecegtgtee aeggaeaaag gggeeateaa 300 gtacaggtge egegaettee teecegaaaa etgeggetge aagateeaet agagaeteat 360 eeaatteeae eatggeegeg egecaeageg geaeageega ttggateett eeatgtteet 420 eegteegtee geettgetae aageaggeag eeaeaeeaa aagetagete tt

- (2) INFORMATION FOR SEQ ID NO:1257:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 89 amino acids

- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..89
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597477
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1257:

Met Lys Ser Ser Thr Leu Leu Ala Ile Leu Val Leu Gln Ala Leu Leu 1 5 10 15

Val Ser Ala Ala Val Ala Lys Asp Leu Gln Gly Arg Arg Arg Ser 20 25 30

Ser Ala Gly Cys Gly Glu Cys Thr Xaa Trp Ser Gly Val Trp Thr Cys 35 40 45

Asp Asp Leu Leu Thr Lys Cys Ala Ala Thr Cys Lys Asn Cys Val Pro 50 60

Val Ser Thr Asp Lys Gly Ala Ile Lys Tyr Arg Cys Arg Asp Phe Leu 65 70 75 80

Pro Glu Asn Cys Gly Cys Lys Ile His 85

- (2) INFORMATION FOR SEQ ID NO:1258:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 406 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..406
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597478
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1258:

 accaaggaaa ttcacaaagc gcctagcaaa agcatacttc ccgaagcact cttgcaatcc 60

 actgagttct gtttgttgag acgcatagag ctagctgcta gcgtcgacaa tgtcgctcgt 120

 gaggcgcacg aacgtgttcg accccttctc gatggacctc tgggatccct tcgacaccat 180

 gttccgctcc atcgtccgt cggcgacctc caccaactcc gagactgccg ccttcgccag 240

 cgcccgcatc gactggaagg agacgccga ggsgcacgtc ttcaaggccg acctcccgg 300

 cgtcaagaag gaggaggtca aggtcgaggt cgaagmggnc aacgtgctgg tcatcagcgg 360

 ccagcgcasa gggagaagga ggacaaggac gacaagtgc accgcg
- (2) INFORMATION FOR SEQ ID NO:1259:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 135 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..135
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597479
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1259:

Thr Lys Glu Ile His Lys Ala Pro Ser Lys Ser Ile Leu Pro Glu Ala 1 5 10 15

Leu Leu Gln Ser Thr Glu Phe Cys Leu Leu Arg Arg Ile Glu Leu Ala 20 25 30

Ala Ser Val Asp Asn Val Ala Arg Glu Ala His Glu Arg Val Arg Pro 35 40 45

Leu Leu Asp Gly Pro Leu Gly Ser Leu Arg His His Val Pro Leu His 50 55 60

Arg Pro Val Gly Asp Leu His Gln Leu Arg Asp Cys Arg Leu Arg Gln

65 70 75 80
Arg Pro His Arg Leu Glu Gly Asp Ala Arg Xaa Ala Arg Leu Gln Gly 85 90 95

Arg Pro Pro Arg Arg Gln Glu Gly Gly Gly Gln Gly Arg Gly Arg Xaa 100 105 110

Xaa Gln Arg Ala Gly His Gln Arg Pro Ala Xaa Arg Glu Lys Glu Asp 115 120 125

Lys Asp Asp Lys Trp His Arg 130 135

- (2) INFORMATION FOR SEQ ID NO:1260:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 99 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..99
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597480
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1260:

Met Ser Leu Val Arg Arg Thr Asn Val Phe Asp Pro Phe Ser Met Asp 1 10 15

Leu Trp Asp Pro Phe Asp Thr Met Phe Arg Ser Ile Val Pro Ser Ala 20 25 30

Thr Ser Thr Asn Ser Glu Thr Ala Ala Phe Ala Ser Ala Arg Ile Asp 35 40 45

Trp Lys Glu Thr Pro Glu Xaa His Val Phe Lys Ala Asp Leu Pro Gly 50 60

Val Lys Lys Glu Glu Val Lys Val Glu Val Glu Xaa Xaa Asn Val Leu 65 70 75 80

Val Ile Ser Gly Gln Arg Xaa Gly Arg Arg Arg Thr Arg Thr Thr Ser 85 90 95

Gly Thr Ala

- (2) INFORMATION FOR SEQ ID NO:1261:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 85 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..85
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597481

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1261:

Ser Ala Thr Ser Thr Asn Ser Glu Thr Ala Ala Phe Ala Ser Ala Arg
20 25 30

Ile Asp Trp Lys Glu Thr Pro Glu Xaa His Val Phe Lys Ala Asp Leu 35 40 45

Pro Gly Val Lys Lys Glu Glu Val Lys Val Glu Val Glu Xaa Xaa Asn 50 55 60

Val Leu Val Ile Ser Gly Gln Arg Xaa Gly Arg Arg Arg Thr Arg Thr 65 70 75 80

Thr Ser Gly Thr Ala

85

- (2) INFORMATION FOR SEQ ID NO:1262:
 - (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 463 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..463
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597482
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1262:

accacageat cegerttee cegaceae egtteetee ageceaggag cacceegeg 60 ceaggaggaa gtttgteet teacaatat gtegagggag gagaatgtet acatggeeaa 120 gttggetgag caggetgaaa ggtatgagga gatggttgag tatatggaga aggtggetaa 240 gaatgtgatt ggggetegee gegetteatg gegeattget tetteeattg aacaaaagga ggagteegt aagaaegaag ageatgtgaa eettateaag gaatateegt gegaagatega 360 ggetgaaetg ageatateet gtgatggaat ettgaagetg ettgaeteec aectagtgee 420 tetteeteet getgetgaat caaaggteet etaceteaag atg

- (2) INFORMATION FOR SEQ ID NO:1263:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 154 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..154
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597483
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1263:

Pro Gln His Pro Xaa Phe Pro Asp His Thr Val Pro Pro Ser Pro Gly
1 10 15

Ala Pro Pro Arg Gln Glu Glu Val Cys Ser Phe Asn Asn Met Ser Arg 20 25 30

Glu Glu Asn Val Tyr Met Ala Lys Leu Ala Glu Gln Ala Glu Arg Tyr 35 40 45

Glu Glu Met Val Glu Tyr Met Glu Lys Val Ala Lys Thr Val Asp Val
50 55 60

Glu Glu Leu Thr Val Glu Glu Arg Asn Leu Leu Ser Val Ala Tyr Lys 65 70 75 80
Asn Val Ile Gly Ala Arg Arg Ala Ser Trp Arg Ile Val Ser Ser Ile

85 90 95
Glu Gln Lys Glu Glu Ser Arg Lys Asn Glu Glu His Val Asn Leu Ile

100 105 110

Lys Glu Tyr Arg Gly Lys Ile Glu Ala Glu Leu Ser Asn Ile Cys Asp

115 120 125

Gly Ile Leu Lys Leu Leu Asp Ser His Leu Val Pro Ser Ser Thr Ala
130 135 140

Ala Glu Ser Lys Val Phe Tyr Leu Lys Met 145 150

- (2) INFORMATION FOR SEQ ID NO:1264:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 125 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..125
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597484

 (xi)
 SEQUENCE DESCRIPTION:
 SEQ ID NO:1264:

 Met
 Ser
 Arg
 Glu
 Glu
 Asn
 Val
 Tyr
 Met
 Ala
 Lys
 Leu
 Ala
 Glu
 Gln
 Ala

 1
 5
 10
 15
 10
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 15
 16
 15
 15
 15
 16
 15
 16
 15
 16
 15
 16
 15
 16
 16
 15
 16
 16
 15
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16
 16<

Asn Leu Ile Lys Glu Tyr Arg Gly Lys Ile Glu Ala Glu Leu Ser Asn 85 90 95

Ile Cys Asp Gly Ile Leu Lys Leu Leu Asp Ser His Leu Val Pro Ser 100 105 110

Ser Thr Ala Ala Glu Ser Lys Val Phe Tyr Leu Lys Met 115 120 125

- (2) INFORMATION FOR SEQ ID NO:1265:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 117 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..117
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597485
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1265:

Met Ala Lys Leu Ala Glu Gln Ala Glu Arg Tyr Glu Glu Met Val Glu

5 10 15

Tyr Met Glu Lys Val Ala Lys Thr Val Asp Val Glu Glu Leu Thr Val 20 25 30

Glu Glu Arg Asn Leu Leu Ser Val Ala Tyr Lys Asn Val Ile Gly Ala 35 40 45

Ser Arg Lys Asn Glu Glu His Val Asn Leu Ile Lys Glu Tyr Arg Gly 65 70 75 80

Lys Ile Glu Ala Glu Leu Ser Asn Ile Cys Asp Gly Ile Leu Lys Leu 85 90 95

Leu Asp Ser His Leu Val Pro Ser Ser Thr Ala Ala Glu Ser Lys Val 100 105 110

Phe Tyr Leu Lys Met 115

- (2) INFORMATION FOR SEQ ID NO:1266:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 401 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..401
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597499
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1266:

agaaacccta acactcatca cgccgccact gatctgctct gttgctggtg actaggagca 60 gtagcgrnga gggattctgg atggccaagt cgaagaacca cacggcgcat aaccagtcgt 120 tcaaggcgca caagaacggc atcaagaagc ccaagcgcca ccgccagacc tccaccaagg ggatggaccc gaagtttctg aggaacttga ggtactcaar gaagggcaac aagaagagtg 240

Page 770

gcgaggctga gtccgaggag taagatgatg gtgattgagt ttgtggcgac aaatgcatct 300 cttgtatcgc taagagtctt agctgtgaac ttagttctca cttctgtgtg actgtgatca 360 aaggacttgt gtatgatcaa aggagatcca tcagttggta c

- (2) INFORMATION FOR SEQ ID NO:1267:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 60 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..60
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597500
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1267:

Met Ala Lys Ser Lys Asn His Thr Ala His Asn Gln Ser Phe Lys Ala 10

His Lys Asn Gly Ile Lys Lys Pro Lys Arg His Arg Gln Thr Ser Thr 25 20

Lys Gly Met Asp Pro Lys Phe Leu Arg Asn Leu Arg Tyr Ser Xaa Lys 40 45

- Gly Asn Lys Lys Ser Gly Glu Ala Glu Ser Glu Glu 50
- (2) INFORMATION FOR SEQ ID NO:1268:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 456 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..456
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597502
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1268:

aaactacacc acaccgcacc ggcaccacca ccaaccgagc ggnnccaggc teetgetcag 60 gaaggggaga agaggcgagc cttccttggg aagtcgcagg aggagagaag gggaacaaag 120 180 atggggcgcg gcaagatcga gatcaagcgg atcgagaact ccaccaaccg ccagtgacct 240 tctccaagcg ccgcaacggg atcctcaaga aggcgcggga gatcagcgtg ctctgcgacg 300 ccgaggtcgg cgtctcgtct tctccagcgc cggcaaagtg taccgggggt cattggtcga 360 gtgcgaggat aactggaatt gccagctcga gaacataacc ttcactgcca aggatgggaa 420 ggtttcgcag ctggagcacg tcttcatcag aggaagcaga gtbagattta tgattatacc tgatatgctc aagaacgctc ccatgttcaa gcgctt

- (2) INFORMATION FOR SEQ ID NO:1269:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 151 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..151
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597503
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1269:

Asn Tyr Thr Thr Pro His Arg His His His Gln Pro Ser Xaa Xaa Arg 10

Leu Leu Leu Arg Lys Gly Arg Arg Gly Glu Pro Ser Leu Gly Ser Arg 25

Arg Arg Arg Glu Gly Glu Gln Arg Trp Gly Ala Ala Arg Ser Arg Ser 45 35 40

Ser Gly Ser Arg Thr Pro Pro Thr Ala Ser Asp Leu Leu Gln Ala Pro 55 Gln Arg Asp Pro Gln Glu Gly Ala Gly Asp Gln Arg Ala Leu Arg Arg 70 75 Arg Gly Arg Arg Leu Val Phe Ser Ser Ala Gly Lys Val Tyr Arg Gly 90 8.5 Ser Leu Val Glu Cys Glu Asp Asn Trp Asn Cys Gln Leu Glu Asn Ile 105 100 Thr Phe Thr Ala Lys Asp Gly Lys Val Ser Gln Leu Glu His Val Phe 125 120 Ile Arg Gly Ser Arg Xaa Arg Phe Met Ile Ile Pro Asp Met Leu Lys 140 135 Asn Ala Pro Met Phe Lys Arg 150 145 (2) INFORMATION FOR SEQ ID NO:1270: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 111 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..111 (D) OTHER INFORMATION: / Ceres Seq. ID 1597504 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1270: Thr Thr Pro His Arg Thr Gly Thr Thr Thr Asn Arg Ala Xaa Pro Gly 10 5 Ser Cys Ser Gly Arg Gly Glu Glu Ala Ser Leu Pro Trp Glu Val Ala 20 25 Gly Gly Glu Lys Gly Asn Lys Asp Gly Ala Arg Gln Asp Arg Asp Gln 45 40 Ala Asp Arg Glu Leu His Gln Pro Pro Val Thr Phe Ser Lys Arg Arg 60 55 Asn Gly Ile Leu Lys Lys Ala Arg Glu Ile Ser Val Leu Cys Asp Ala 75 70 Glu Val Gly Val Ser Ser Pro Ala Pro Ala Lys Cys Thr Gly Gly 90 85 His Trp Ser Ser Ala Arg Ile Thr Gly Ile Ala Ser Ser Arg Thr 105 100 (2) INFORMATION FOR SEQ ID NO:1271: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 494 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: -(B) LOCATION: 1..494 (D) OTHER INFORMATION: / Ceres Seq. ID 1597519

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1271:	
atcaactcgc aattccgcat tcgttccaac tcctgagcac tttctcctcg tcccagacaa	60
gccaatcgaa gccgagatac gaccatggac tccaccggca ccggagcagg gggtaagggg	120
aagaagggag cggccgggcg caaggtcggc gggccaagga agaagtcggt gtcgaggtcc	180
gtgaaggccg gtctccagtt ccccgtcggc cgcatcgggc gctacttgaa gaagggccgc	240
tacgogoang ttoggoacog gogococogt ctacctogoc gotgtoctog aatacctogo	300
cgctgaggtt ctggagctcg ccggtaatgc tgccagggac aaggcgatca agaggtttca	360
ggtkaggaac atcgtggaac aggcggccat cagggacgtc caggaagcat gcgtctatga	420
tggctacgtc ctcccgaagc tgtacgccaa ggtgcaccac tgcgtctcgt gcgcgatcca	480
mgstcacatc gtcc	

- Attorney Docket No 750-1237P 46.003 Client Docket No. (2) INFORMATION FOR SEQ ID NO:1272: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 60 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..60 (D) OTHER INFORMATION: / Ceres Seq. ID 1597520 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1272: Ser Thr Arg Asn Ser Ala Phe Val Pro Thr Pro Glu His Phe Leu Leu 10 Val Pro Asp Lys Pro Ile Glu Ala Glu Ile Arg Pro Trp Thr Pro Pro 30 20 25 Ala Pro Glu Gln Gly Val Arg Gly Arg Arg Glu Arg Pro Gly Ala Arg 40 Ser Ala Gly Gln Gly Arg Ser Arg Cys Arg Gly Pro 55 50 (2) INFORMATION FOR SEQ ID NO:1273: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 73 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1...73 (D) OTHER INFORMATION: / Ceres Seq. ID 1597521 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1273: Met Asp Ser Thr Gly Thr Gly Ala Gly Gly Lys Gly Lys Gly Ala 5 10 Ala Gly Arg Lys Val Gly Gly Pro Arg Lys Lys Ser Val Ser Arg Ser 25 Val Lys Ala Gly Leu Gln Phe Pro Val Gly Arg Ile Gly Arg Tyr Leu 40 4.5 Lys Lys Gly Arg Tyr Ala Xaa Val Arg His Arg Arg Pro Arg Leu Pro - 55 60 Arg Arg Cys Pro Arg Ile Pro Arg Arg 70 (2) INFORMATION FOR SEQ ID NO:1274: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 56 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..56
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597522

 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1274:

 Met Leu Pro Gly Thr Arg Arg Ser Arg Gly Phe Arg Xaa Gly Thr Ser

 1 5 10 15

 Trp Asn Arg Arg Pro Ser Gly Thr Ser Arg Lys His Ala Ser Met Met
 20 25 30

 Ala Thr Ser Ser Arg Ser Cys Thr Pro Arg Cys Thr Thr Ala Ser Arg
 35 40 45

 Ala Arg Ser Xaa Xaa Thr Ser Ser

₹50-1237P

50 55

- (2) INFORMATION FOR SEQ ID NO:1275:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 364 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..364
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597540
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1275:

aaactcgcaa ttccgcattc gttccaactc ctgagcactt tctcctcgtc cccgacaagc 60 caatcgaagc cgagatacgg ccatggactc caccggcacc ggagcagggg gtaaggggaa 120 gangggagcg gccgggcgca aggtcggcgg gccaaggaag aagtcggtgt cgaggtccgt 180 gaaggccggg ctccagttcc ccgtcggccg catcgggcgc tacttgaaga agggccgcta 240 egegeastgt neggeaeegg egeeeeegte taeetegemg etgteetega ataeetekee 300 gkaatttctg mcaactmntd gtaatsctgc sasrgacamc agaagactcg cattattccr 360 cacc

- (2) INFORMATION FOR SEQ ID NO:1276:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 59 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..59
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597541
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1276:

Thr Arg Asn Ser Ala Phe Val Pro Thr Pro Glu His Phe Leu Leu Val

Pro Asp Lys Pro Ile Glu Ala Glu Ile Arg Pro Trp Thr Pro Pro Ala 20 25

Pro Glu Gln Gly Val Arg Gly Arg Xaa Glu Arg Pro Gly Ala Arg Ser 40

Ala Gly Gln Gly Arg Ser Arg Cys Arg Gly Pro

- (2) INFORMATION FOR SEQ ID NO:1277:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 94 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..94
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597542
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1277:

Met Asp Ser Thr Gly Thr Gly Ala Gly Gly Lys Gly Lys Xaa Gly Ala 10 1.5

Ala Gly Arg Lys Val Gly Gly Pro Arg Lys Lys Ser Val Ser Arg Ser 25 20

Val Lys Ala Gly Leu Gln Phe Pro Val Gly Arg Ile Gly Arg Tyr Leu 40 45

Lys Lys Gly Arg Tyr Ala Xaa Cys Xaa Ala Pro Ala Pro Pro Ser Thr 55 50 60

Ser Xaa Leu Ser Ser Asn Thr Xaa Pro Xaa Phe Leu Xaa Thr Xaa Xaa

65 70 75 80

Asn Xaa Xaa Xaa Asp Xaa Arg Arg Leu Ala Leu Phe Xaa Ala 85 90

- (2) INFORMATION FOR SEQ ID NO:1278:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 479 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..479
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597553
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1278: attttcggca agattgcaac agggtcgtcc ggtcgcgttt ctcctacgcg gtccgtgggc tgtgctgcct gctgttgcat cgtctcctcg tttaaacggg cgaactcaag attgcaacag 120 ggtcgtccgc cgccaccaat tgctagcgca atcctagccc tagcccagcg ccctccctg 180 cgtcgtgctc ggtagcgatg gcgtcgtttt ctgaagcgcc cccggggaac ccaactgctg 240 gcgagaagat cttcaagacc aagtgcgcgc agtgccacac cgtcgacaag ggcgccggcc 300 acaagcaagg toctaatttg aatggtotgt ttgggaggca gtotggtaca acacotggot 360 actcttactc ttcggctaac aagaacatgg ctgtgatttg ggaggagaac actttgtatg 420 actacttgct taatcccaag aagtacattc ctggaaccaa gatggtattc cctggactg (2) INFORMATION FOR SEQ ID NO:1279:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 64 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..64
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597554
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1279:
 Ile Phe Gly Lys Ile Ala Thr Gly Ser Ser Gly Arg Val Ser Pro Thr

1 5 10 15
Arg Ser Val Gly Cys Ala Ala Cys Cys Cys Ile Val Ser Ser Phe Lys

20 25 30

Arg Ala Asn Ser Arg Leu Gln Gln Gly Arg Pro Pro Pro Ile Ala
35 40 45

Ser Ala Ile Leu Ala Leu Ala Gln Arg Pro Ser Leu Arg Arg Ala Arg

- (2) INFORMATION FOR SEQ ID NO:1280:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 51 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..51
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597555

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1280:

Phe Ser Ala Arg Leu Gln Gln Gly Arg Pro Val Ala Phe Leu Arg
1 5 10 15

Gly Pro Trp Ala Val Leu Pro Ala Val Ala Ser Ser Pro Arg Leu Asn 20 25 30

```
Page 775
Client Docket No.
Gly Arg Thr Gln Asp Cys Asn Arg Val Val Arg Arg His Gln Leu Leu
                            40
Ala Gln Ser
    50
(2) INFORMATION FOR SEQ ID NO:1281:
     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 94 amino acids
          (B) TYPE: amino acid
          (C) STRANDEDNESS:
          (D) TOPOLOGY: linear
    (ii) MOLECULE TYPE: peptide
    (ix) FEATURE:
          (A) NAME/KEY: peptide
          (B) LOCATION: 1..94
          (D) OTHER INFORMATION: / Ceres Seq. ID 1597556
    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1281:
Met Ala Ser Phe Ser Glu Ala Pro Pro Gly Asn Pro Thr Ala Gly Glu
                                     10
Lys Ile Phe Lys Thr Lys Cys Ala Gln Cys His Thr Val Asp Lys Gly
                                                     30
                                 25
Ala Gly His Lys Gln Gly Pro Asn Leu Asn Gly Leu Phe Gly Arg Gln
                                                 45
                             40
Ser Gly Thr Thr Pro Gly Tyr Ser Tyr Ser Ser Ala Asn Lys Asn Met
                         55
                                             60
Ala Val Ile Trp Glu Glu Asn Thr Leu Tyr Asp Tyr Leu Leu Asn Pro
                     70
                                         75
Lys Lys Tyr Ile Pro Gly Thr Lys Met Val Phe Pro Gly Leu
                                     90
                85
(2) INFORMATION FOR SEQ ID NO:1282:
     (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 116 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear
     (ii) MOLECULE TYPE: DNA (genomic)
     (ix) FEATURE:
           (A) NAME/KEY: -
           (B) LOCATION: 1..116
           (D) OTHER INFORMATION: / Ceres Seq. ID 1597579
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1282:
cnaaaaaagg tacacatcat ctcttccaaa tcccaccagc aaatccccaa ttcctccgag
                                                                         60
 ttcgagcgac catgtcgggg cgtggcaagg gcgggaaggg tctgggcaaa ggcggc
 (2) INFORMATION FOR SEQ ID NO:1283:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 38 amino acids
           (B) TYPE: amino acid
           (C) STRANDEDNESS:
           (D) TOPOLOGY: linear
     (ii) MOLECULE TYPE: peptide
     (ix) FEATURE:
           (A) NAME/KEY: peptide
           (B) LOCATION: 1..38
           (D) OTHER INFORMATION: / Ceres Seq. ID 1597580
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1283:
 Xaa Lys Lys Val His Ile Ile Ser Ser Lys Ser His Gln Gln Ile Pro
                                     10
                 5
 Asn Ser Ser Glu Phe Glu Arg Pro Cys Arg Gly Val Ala Arg Ala Gly
                                  25
             20
```

(2) INFORMATION FOR SEQ ID NO:1284:

Arg Val Trp Ala Lys Ala

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 38 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..38
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597581
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1284:

Xaa Lys Arg Tyr Thr Ser Ser Leu Pro Asn Pro Thr Ser Lys Ser Pro 1 5 10 15

Ile Pro Pro Ser Ser Ser Asp His Val Gly Ala Trp Gln Gly Arg Glu 20 25 30

Gly Ser Gly Gln Arg Arg 35

- (2) INFORMATION FOR SEQ ID NO:1285:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 38 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..38
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597582
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1285:

Lys Lys Gly Thr His His Leu Phe Gln Ile Pro Pro Ala Asn Pro Gln 1 5 10 15

Phe Leu Arg Val Arg Ala Thr Met Ser Gly Arg Gly Lys Gly Gly Lys 20 25 30

Gly Leu Gly Lys Gly Gly 35

- (2) INFORMATION FOR SEQ ID NO:1286:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 486 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..486
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597589

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1286:

accgcagcca caccccaaca tcaaaaccct cccatctctt cccctccgcc gccgtctctg 60 ctctcaagat cgccatggcg tcagaaaaga agcagtcgaa cccgatgcgg gagatcaagg 120 tgcagaagct cgtgctcaat atctccgttg gagagagtgg agatcgactc acccgcgccg 180 ccaaggtgtt ggagcagttg agtgggcagt ctccggtttt ctccaaggca aggtatactg 240 tgaggtcttt cggtattcgt cgtaatgaga agattgcatg ctacgttacg gtcaggggtg 300 agaaggcaat gcagcttctt gagagtggcc tcaaggtgaa ggagtatgag cttctgagga 360 ggaacttcag tgatactggc tgctttggtt ttggtattca ggagcacatt gatcttggaa 420 tcaagtatga cccgtctact ggtatttatn gcatggactt ttacgtcgtt ctggaacgag 480 ctcqtt

- (2) INFORMATION FOR SEQ ID NO:1287:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 161 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..161
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597590
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1287:
- Arg Ser His Thr Pro Thr Ser Lys Pro Ser His Leu Phe Pro Ser Ala 1 5 10 15
- Ala Val Ser Ala Leu Lys Ile Ala Met Ala Ser Glu Lys Lys Gln Ser 20 25 30
- Asn Pro Met Arg Glu Ile Lys Val Gln Lys Leu Val Leu Asn Ile Ser 35 40 45
- Val Gly Glu Ser Gly Asp Arg Leu Thr Arg Ala Ala Lys Val Leu Glu 50 60
- Gln Leu Ser Gly Gln Ser Pro Val Phe Ser Lys Ala Arg Tyr Thr Val 65 70 75 80
- Arg Ser Phe Gly Ile Arg Arg Asn Glu Lys Ile Ala Cys Tyr Val Thr 85 90 95
- Val Arg Gly Glu Lys Ala Met Gln Leu Leu Glu Ser Gly Leu Lys Val 100 105 110
- Lys Glu Tyr Glu Leu Leu Arg Arg Asn Phe Ser Asp Thr Gly Cys Phe 115 120 125
- Gly Phe Gly Ile Gln Glu His Ile Asp Leu Gly Ile Lys Tyr Asp Pro 130 135 140
- Ser Thr Gly Ile Tyr Xaa Met Asp Phe Tyr Val Val Leu Glu Arg Ala 145 150 155 160 Arg
- (2) INFORMATION FOR SEQ ID NO:1288:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 137 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..137
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597591
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1288:
- Met Ala Ser Glu Lys Lys Gln Ser Asn Pro Met Arg Glu Ile Lys Val

 1 10 15
- Gln Lys Leu Val Leu Asn Ile Ser Val Gly Glu Ser Gly Asp Arg Leu 20 25 30
- Thr Arg Ala Ala Lys Val Leu Glu Gln Leu Ser Gly Gln Ser Pro Val 35 40 45
- Phe Ser Lys Ala Arg Tyr Thr Val Arg Ser Phe Gly Ile Arg Arg Asn 50 55 60
- Glu Lys Ile Ala Cys Tyr Val Thr Val Arg Gly Glu Lys Ala Met Gln 65 70 75 80
- Leu Leu Glu Ser Gly Leu Lys Val Lys Glu Tyr Glu Leu Leu Arg Arg 85 90 95
- Asn Phe Ser Asp Thr Gly Cys Phe Gly Phe Gly Ile Gln Glu His Ile 100 105 110
- Asp Leu Gly Ile Lys Tyr Asp Pro Ser Thr Gly Ile Tyr Xaa Met Asp 115 120 125
- Phe Tyr Val Val Leu Glu Arg Ala Arg 130 135
- (2) INFORMATION FOR SEQ ID NO:1289:
 - (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 127 amino acids(B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..127
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597592
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1289:
- Met Arg Glu Ile Lys Val Gln Lys Leu Val Leu Asn Ile Ser Val Gly
 1 10 15
- Glu Ser Gly Asp Arg Leu Thr Arg Ala Ala Lys Val Leu Glu Gln Leu 20 25 30
- Ser Gly Gln Ser Pro Val Phe Ser Lys Ala Arg Tyr Thr Val Arg Ser 35 40 45
- Phe Gly Ile Arg Arg Asn Glu Lys Ile Ala Cys Tyr Val Thr Val Arg 50 55 60
- Gly Glu Lys Ala Met Gln Leu Glu Ser Gly Leu Lys Val Lys Glu
 65 70 75 80
- Tyr Glu Leu Leu Arg Arg Asn Phe Ser Asp Thr Gly Cys Phe Gly Phe 85 90 95
- Gly Ile Gln Glu His Ile Asp Leu Gly Ile Lys Tyr Asp Pro Ser Thr 100 105 110
- Gly Ile Tyr Xaa Met Asp Phe Tyr Val Val Leu Glu Arg Ala Arg 115 120 125
- (2) INFORMATION FOR SEQ ID NO:1290:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 413 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..413
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597600
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1290:
- accgacggca tttgcatctk agttcctgat tgttgtattc cagtttcttc tgtgagtttt 60 gtgggatcgc gaggwagaac tagaaggatg tcgtgctgcg gaggcaactg cgggtkcggc 120 gccggctgca agtgcggcac ggctgcggag ggtgcaagat gtacccggac atggctgagc aggtgaccac cactacccag actctcatca tgggtgttgc accgtccaag ggcgggttcg 240 aggcggcgc gangctgaga acggcggtg caagtkcggc gccaactgca cctgcgaccc ctgcacctgc aagtgagacg acggcgatga cgctkcaggg cgtcaattgg gatcggaagg 360 actcttttat cttaagctct agtataatat tctagcagtg tcgtytgtgt ttt
- (2) INFORMATION FOR SEQ ID NO:1291:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 61 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..61
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597601
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1291:
- Thr Asp Gly Ile Cys Ile Xaa Val Pro Asp Cys Cys Ile Pro Val Ser 1 5 10 15
- Ser Val Ser Phe Val Gly Ser Arg Xaa Arg Thr Arg Arg Met Ser Cys
 20 25 30

Cys Gly Gly Asn Cys Gly Xaa Gly Ala Gly Cys Lys Cys Gly Thr Ala

Ala Glu Gly Ala Arg Cys Thr Arg Thr Trp Leu Ser Arg 50 55 60

- (2) INFORMATION FOR SEQ ID NO:1292:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 78 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..78
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597602
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1292:

Met Tyr Pro Asp Met Ala Glu Gln Val Thr Thr Thr Gln Thr Leu
1 10 15

Ile Met Gly Val Ala Pro Ser Lys Gly Gly Phe Glu Ala Ala Ala Xaa 20 25 30

Leu Arg Thr Ala Gly Ala Ser Xaa Ala Pro Thr Ala Pro Ala Thr Pro 35 40 45

Ala Pro Ala Ser Glu Thr Thr Ala Met Thr Xaa Gln Gly Val Asn Trp 50 60

Asp Arg Lys Asp Ser Phe Ile Leu Ser Ser Ser Ile Ile Phe 65 70 75

- (2) INFORMATION FOR SEQ ID NO:1293:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 74 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..74
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597603
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1293:

Met Ala Glu Gln Val Thr Thr Thr Gln Thr Leu Ile Met Gly Val 1 5 10 15

Ala Pro Ser Lys Gly Gly Phe Glu Ala Ala Ala Xaa Leu Arg Thr Ala 20 25 30

Gly Ala Ser Xaa Ala Pro Thr Ala Pro Ala Thr Pro Ala Pro Ala Ser 35 40 45

Glu Thr Thr Ala Met Thr Xaa Gln Gly Val Asn Trp Asp Arg Lys Asp 50 55 60

Ser Phe Ile Leu Ser Ser Ser Ile Ile Phe 65 70

- (2) INFORMATION FOR SEQ ID NO:1294:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 449 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..449
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597620
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1294:
- gaagagggaa ggggggatat cggggaaaaa aacacaaggt accagcagcg caaagagccg

ggtgttggca cagatcaggc ggccgcatgg cgatcmgttc gcggattcgg cgaacaacgt 120 gatcatcgag gaggtgaaca agggcctgaa cccaggaatg gtggttctgc tcgatggttg 180 ccagcttcct gctgctcttc ttcgccgga actacggct gtatgtgtac gcgcagaaga 240 cgctcccgcc gaagaagaag aagccggtct ccaagaggaa gctgaagagg gaaaagctga 300 agcaggggt ctcggcgca ggggagtgaa cagacaggga tgccgccgat cggcccgtcc 360 tgagcccgcc ccctatttac atgattgtcg aagaaaaaag atacacctgc tggagccgcg 420 gtgttcgtag taattttgtt ctgtagatg

- (2) INFORMATION FOR SEQ ID NO:1295:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 94 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..94
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597621
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1295:

Glu Glu Gly Arg Gly Asp Ile Gly Glu Lys Asn Thr Arg Tyr Gln Gln
1 10 15

Arg Lys Glu Pro Gly Val Gly Thr Asp Gln Ala Ala Ala Trp Arg Xaa 20 25 30

Val Arg Gly Phe Gly Glu Gln Arg Asp His Arg Gly Gly Glu Gln Gly
35 40 45

Pro Glu Pro Arg Asn Gly Gly Ser Ala Arg Trp Leu Pro Ala Ser Cys 50 55 60

Cys Ser Ser Ser Pro Gly Thr Thr Arg Cys Met Cys Thr Arg Arg Arg 65 70 75 80

Arg Ser Arg Arg Arg Arg Ser Arg Ser Pro Arg Gly Ser 85 90

- (2) INFORMATION FOR SEQ ID NO:1296:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 142 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..142
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597622
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1296:

Lys Arg Glu Gly Gly Ile Ser Gly Lys Lys Thr Gln Gly Thr Ser Ser 1 10 15

Ala Lys Ser Arg Val Leu Ala Gln Ile Arg Arg Pro His Gly Asp Xaa 20 25 30

Phe Ala Asp Ser Ala Asn Asn Val Ile Ile Glu Glu Val Asn Lys Gly 35 40 45

Leu Asn Pro Gly Met Val Val Leu Leu Asp Gly Cys Gln Leu Pro Ala 50 55 60

Ala Leu Leu Arg Arg Glu Leu Arg Ala Val Cys Val Arg Ala Glu Asp
65 70 75 80

Ala Pro Ala Glu Glu Glu Ala Gly Leu Gln Glu Glu Ala Glu Glu 85 90 95

Gly Lys Ala Glu Ala Gly Gly Leu Gly Ala Arg Gly Val As
n Arg Gl
n $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110 \hspace{1.5cm}$

Gly Cys Arg Arg Ser Ala Arg Pro Glu Pro Ala Pro Tyr Leu His Asp 115 120 125

Cys Arg Arg Lys Lys Ile His Leu Leu Glu Pro Arg Cys Ser 130 135 140

- Attorney Docket No 750-1237P Client Docket No. ' 46.003 (2) INFORMATION FOR SEQ ID NO:1297: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 90 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..90 (D) OTHER INFORMATION: / Ceres Seq. ID 1597623 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1297: Met Val Val Leu Leu Asp Gly Cys Gln Leu Pro Ala Ala Leu Leu Arg 10 Arg Glu Leu Arg Ala Val Cys Val Arg Ala Glu Asp Ala Pro Ala Glu 20 25 Glu Glu Glu Ala Gly Leu Gln Glu Glu Ala Glu Glu Gly Lys Ala Glu 40
- Ala Gly Gly Leu Gly Ala Arg Gly Val Asn Arg Gln Gly Cys Arg Arg
 50

 Ser Ala Arg Pro Glu Pro Ala Pro Tyr Leu His Asp Cys Arg Arg Lys

65 70 75
Lys Ile His Leu Leu Glu Pro Arg Cys Ser

- 85
 (2) INFORMATION FOR SEQ ID NO:1298:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 385 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..385
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597631
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1298:

 aatcaactcg caattccgca ttcgttccaa ctcctgagca ctttctcctc gtcccagaca agccaatcga agccgagata cgaccatgga ctccaccggc accggagcag ggggtaaggg gaagaaggga gcggccgggc gcaaggtcgg cgggccaagg aagaagtcgg tgtcgaggtc cgtgaaggcc ggtctccagt tccccgtcgg ccgcatcggg cgctacttka agaagggccg ctacggcab gttcggcacc ggcgccccg tctacctcgc cgctgtcctc gaatacctcg cgctgaggt tctggaggt tctggaggt ctgcaggga caacaagaag actcgcatta 360
- (2) INFORMATION FOR SEQ ID NO:1299:

ttccgcgcca cgtgcttctk gcgat

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 38 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..38
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597632
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1299:

Asn Gln Leu Ala Ile Pro His Ser Phe Gln Leu Leu Ser Thr Phe Ser 1 5 5 10 15 Ser Ser Gln Thr Ser Gln Ser Lys Pro Arg Tyr Asp His Gly Leu His

Ser Ser Gin Thr Ser Gin Ser Lys Pro Arg Tyr Asp his Gry Led his

Arg His Arg Ser Arg Gly

```
750-1237P
Client Docket No.
(2) INFORMATION FOR SEQ ID NO:1300:
     (i) SEQUENCE CHARACTERISTICS:
          (A) LENGTH: 60 amino acids
          (B) TYPE: amino acid
          (C) STRANDEDNESS:
          (D) TOPOLOGY: linear
    (ii) MOLECULE TYPE: peptide
    (ix) FEATURE:
          (A) NAME/KEY: peptide
          (B) LOCATION: 1..60
          (D) OTHER INFORMATION: / Ceres Seq. ID 1597633
    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1300:
Ser Thr Arg Asn Ser Ala Phe Val Pro Thr Pro Glu His Phe Leu Leu
                                     10
                5
Val Pro Asp Lys Pro Ile Glu Ala Glu Ile Arg Pro Trp Thr Pro Pro
                                                     30
                                 25
            20
Ala Pro Glu Gln Gly Val Arg Gly Arg Arg Glu Arg Pro Gly Ala Arg
                             40
Ser Ala Gly Gln Gly Arg Ser Arg Cys Arg Gly Pro
                         55
(2) INFORMATION FOR SEQ ID NO:1301:
     (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 73 amino acids
           (B) TYPE: amino acid
           (C) STRANDEDNESS:
           (D) TOPOLOGY: linear
    (ii) MOLECULE TYPE: peptide
     (ix) FEATURE:
           (A) NAME/KEY: peptide
           (B) LOCATION: 1..73
           (D) OTHER INFORMATION: / Ceres Seq. ID 1597634
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1301:
Met Asp Ser Thr Gly Thr Gly Ala Gly Gly Lys Gly Lys Gly Ala
                 5
Ala Gly Arg Lys Val Gly Gly Pro Arg Lys Lys Ser Val Ser Arg Ser
                                                      30
                                 25
             20
Val Lys Ala Gly Leu Gln Phe Pro Val Gly Arg Ile Gly Arg Tyr Xaa
                                                  45
                             40
Lys Lys Gly Arg Tyr Ala Xaa Val Arg His Arg Arg Pro Arg Leu Pro
                                              60
                         55
 Arg Arg Cys Pro Arg Ile Pro Arg Arg
                     70
 (2) INFORMATION FOR SEQ ID NO:1302:
      (i) SEQUENCE CHARACTERISTICS:
           (A) LENGTH: 470 base pairs
           (B) TYPE: nucleic acid
           (C) STRANDEDNESS: single
           (D) TOPOLOGY: linear
     (ii) MOLECULE TYPE: DNA (genomic)
     (ix) FEATURE:
           (A) NAME/KEY: -
            (B) LOCATION: 1..470
           (D) OTHER INFORMATION: / Ceres Seq. ID 1597639
     (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1302:
```

aacaatgcca attgccagca acccatccat actttcagct gttgatacaa aaagagaaga 60 gagaatggcg gavaacccag agttgagtgg tcttattact caaacaaaga agaaaaatat 120 agtgccaaag ttcaccgaaa tcttccccgt ggaggacacg cctaccctta cagcgccttc 180 atcacctccg tccggaaaga agtgatcaaa tactgcacca accatacagg catcgtccag 240 cccgtgctgc cgctggagaa gaatgtcccc gagctctggt tctacaccga gctcaaaacg 300 aagaccaggt ccatcacgct cgccatacgt atggacaacc tctacctggt cggcttcagg 360 acccccggcg gtgtggtggg agttcggcaa ggacggcgac acccacctcc tcgacgacaa 420

46.003 Client Docket No. cgccaagtgg ctcggctttg gcgggccggt accaggacct catcgggcag (2) INFORMATION FOR SEQ ID NO:1303: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 67 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..67 (D) OTHER INFORMATION: / Ceres Seq. ID 1597640 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1303: Thr Met Pro Ile Ala Ser Asn Pro Ser Ile Leu Ser Ala Val Asp Thr 10 5 Lys Arg Glu Glu Arg Met Ala Xaa Asn Pro Glu Leu Ser Gly Leu Ile 30 25 20 Thr Gln Thr Lys Lys Lys Asn Ile Val Pro Lys Phe Thr Glu Ile Phe 45 40 Pro Val Glu Asp Thr Pro Thr Leu Thr Ala Pro Ser Ser Pro Pro Ser 55 Gly Lys Lys 65 (2) INFORMATION FOR SEQ ID NO:1304: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 66 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..66 (D) OTHER INFORMATION: / Ceres Seq. ID 1597641 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1304: Met Pro Ile Ala Ser Asn Pro Ser Ile Leu Ser Ala Val Asp Thr Lys 10 5 Arg Glu Glu Arg Met Ala Xaa Asn Pro Glu Leu Ser Gly Leu Ile Thr 25 Gln Thr Lys Lys Lys Asn Ile Val Pro Lys Phe Thr Glu Ile Phe Pro 45 40 Val Glu Asp Thr Pro Thr Leu Thr Ala Pro Ser Ser Pro Pro Ser Gly 60 50 55 Lys Lys 65 (2) INFORMATION FOR SEQ ID NO:1305: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 69 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..69 (D) OTHER INFORMATION: / Ceres Seq. ID 1597642 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1305: Met Ser Pro Ser Ser Gly Ser Thr Pro Ser Ser Lys Arg Arg Pro Gly

10

Pro Ser Arg Ser Pro Tyr Val Trp Thr Thr Ser Thr Trp Ser Ala Ser 25

Gly Pro Pro Ala Val Trp Trp Glu Phe Gly Lys Asp Gly Asp Thr His

Leu Leu Asp Asp Asn Ala Lys Trp Leu Gly Phe Gly Gly Pro Val Pro 50 55 60

Gly Pro His Arg Ala

65

- (2) INFORMATION FOR SEQ ID NO:1306:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 371 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..371
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597658
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1306:

 accaaggaaa ttcacaaagc gatactagct agtccctacc aaagcatact tcctgaaaca 60
 ctcttgcaat ccactgagtt ctgtttgttg agacgcatag agctagagtc gacaatgtcg
 ctcgtgaggc gcasaacgtg ttcgacccct tctccatgga cctctgggac cccttcgaca 180
 ccatgttccg ctccatcgtc ccgtcggcga cctccaccaa ctccgagact gccgccttcg
 ccagcgcccg catcgactgg aaggagacgc ccgaggcga cgtgttcaag gccgacctcc
 ccgqcqtcaa gaaggaagag gtcaaggtcg aggtcgagga cggcaacatg ctggtcatca 360
- (2) INFORMATION FOR SEQ ID NO:1307:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 123 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

gcggccagcg c

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..123
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597659
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1307:

Thr Lys Glu Ile His Lys Ala Ile Leu Ala Ser Pro Tyr Gln Ser Ile 1 5 10 15

Leu Pro Glu Thr Leu Leu Gln Ser Thr Glu Phe Cys Leu Leu Arg Arg 20 25 30

Ile Glu Leu Glu Ser Thr Met Ser Leu Val Arg Arg Xaa Thr Cys Ser 35 40 45

Thr Pro Ser Pro Trp Thr Ser Gly Thr Pro Ser Thr Pro Cys Ser Ala 50 55 60

Pro Ser Ser Arg Arg Pro Pro Pro Thr Pro Arg Leu Pro Pro Ser 65 70 75 80

Pro Ala Pro Ala Ser Thr Gly Arg Arg Pro Arg Arg Thr Cys Ser 85 90 95

Arg Pro Thr Ser Pro Ala Ser Arg Arg Lys Arg Ser Arg Ser Arg Ser 100 105 110

Arg Thr Ala Thr Cys Trp Ser Ser Ala Ala Ser 115 120

- (2) INFORMATION FOR SEQ ID NO:1308:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 85 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..85
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597660
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1308:
- Met Ser Leu Val Arg Arg Xaa Thr Cys Ser Thr Pro Ser Pro Trp Thr

 1 10 15

 Can Clu The Dec Can The Dec Can Ala Dec Can Arg Arg Arg
- Ser Gly Thr Pro Ser Thr Pro Cys Ser Ala Pro Ser Ser Arg Arg Arg 20 25 30
- Pro Pro Pro Thr Pro Arg Leu Pro Pro Ser Pro Ala Pro Ala Ser Thr
 35 40 45
- Gly Arg Arg Pro Arg Arg Thr Cys Ser Arg Pro Thr Ser Pro Ala 50 55 60
- Ser Arg Arg Lys Arg Ser Arg Ser Arg Ser Arg Thr Ala Thr Cys Trp 65 70 75 80
- Ser Ser Ala Ala Ser 85
- (2) INFORMATION FOR SEQ ID NO:1309:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 72 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..72
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597661
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1309:
- Met Asp Leu Trp Asp Pro Phe Asp Thr Met Phe Arg Ser Ile Val Pro
 1 10 15
- Ser Ala Thr Ser Thr Asn Ser Glu Thr Ala Ala Phe Ala Ser Ala Arg 20 25 30
- Ile Asp Trp Lys Glu Thr Pro Glu Ala His Val Phe Lys Ala Asp Leu 35 40 45
- Pro Gly Val Lys Lys Glu Glu Val Lys Val Glu Val Glu Asp Gly Asn
- Met Leu Val Ile Ser Gly Gln Arg 65 70
- (2) INFORMATION FOR SEQ ID NO:1310:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 236 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..236
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597666
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1310:
- acgcggaaac cctagcagct caacacctc gaggtacctc cattcgcagt cggccgtgtc 60 ctcggagccg cgggagatgg cggatcagga gaccgtagct gtggttgagg caccaacccc ggttcttggg gaacgatgga cctgatgact gcctgcarc tcgtcatgaa gaagtcaggt 180 gctcacgatk gccttttgaa gggtcttcgt gaggctgcca aagcatcgag aagcat
- (2) INFORMATION FOR SEQ ID NO:1311:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 47 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide

(ix) FEATURE:

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..47
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597667

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1311:

Arg Gly Asn Pro Ser Ser Ser Thr Pro Ser Arg Tyr Leu His Ser Gln 1 10 15

Ser Ala Val Ser Ser Glu Pro Arg Glu Met Ala Asp Gln Glu Thr Val 20 25 30

Ala Val Val Glu Ala Pro Thr Pro Val Leu Gly Glu Arg Trp Thr 35 40 45

- (2) INFORMATION FOR SEQ ID NO:1312:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 31 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..31
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597668
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1312:

Ala Glu Thr Leu Ala Ala Gln His Pro Arg Gly Thr Ser Ile Arg Ser

1 10 15

Arg Pro Cys Pro Arg Ser Arg Gly Arg Trp Arg Ile Arg Arg Pro 20 25 30

- (2) INFORMATION FOR SEQ ID NO:1313:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..33
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597669

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1313:

Met Asp Leu Met Thr Ala Leu Xaa Leu Val Met Lys Lys Ser Gly Ala 1 5 10 15

His Asp Xaa Leu Leu Lys Gly Leu Arg Glu Ala Ala Lys Ala Ser Arg 20 25 30

Ser

- (2) INFORMATION FOR SEQ ID NO:1314:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 401 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..401
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597674
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1314:

ttctgaagtt tcggcggcgg cgtaggtagc ggcggtgtgc ttgtgatctc aatctcaacc 60 ccaggcgagc tcggctccgg tcacccgtcg atccacgcaa ccatgtcgag gaggaagacc 120 agggagccca aggaggagaa cgttaccctt ggacccactg tccgtgaagg agagtttgtb tttggtgttg ctcacatctt tgcatccttc aatgacacct tcattcatgt cactgatttg 240

46.003

tctgggaggg aaactttggt tcggatcact ggtggcatga aggttaaggc tgatcgtgat 300 gagtcgtctc cttatgctgc tatgcttgct gcccaagatg ttgcacagcg ttgcaaggag 360 ctcggtatca cagcgctgca cattaagctt cgtgccactg g

- (2) INFORMATION FOR SEQ ID NO:1315:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 99 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..99
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597675
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1315:

Met Ser Arg Arg Lys Thr Arg Glu Pro Lys Glu Glu Asn Val Thr Leu 10

Gly Pro Thr Val Arg Glu Gly Glu Phe Xaa Phe Gly Val Ala His Ile 25 20

Phe Ala Ser Phe Asn Asp Thr Phe Ile His Val Thr Asp Leu Ser Gly 40

Arg Glu Thr Leu Val Arg Ile Thr Gly Gly Met Lys Val Lys Ala Asp 55 60 50

Arg Asp Glu Ser Ser Pro Tyr Ala Ala Met Leu Ala Ala Gln Asp Val 70

Ala Gln Arg Cys Lys Glu Leu Gly Ile Thr Ala Leu His Ile Lys Leu 90 85

Arg Ala Thr

- (2) INFORMATION FOR SEQ ID NO:1316:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 399 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..399
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597698
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1316:

catcaatagc cagttctaat ctcaggttrg gggaaatagg cctcgcgggt cgcaaaccgc 60 120 tctctgctga tttrggattt cggaggcgct cgagttttca ggtaaactat agtttagtga cccatggctc gtactaagca aactgctcgc aagtccactg gagggaaggc tccwaggaaa 180 caacttgcca ccaaggctgc ccgtaagtct gcacctacaa ctgggggagt aaagaagcct 240 caccgttacc gccctggaac tgttgctctt cgtgaaatcc gcaagtatca gaagagcact 300 gacctgctca taaggaagct tccgttccaa aggcttgtca gggagattkc ccaggatttc 360 aagactgatc ttcgtttcca gagccatgcg gtgcttgcg

- (2) INFORMATION FOR SEQ ID NO:1317:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 92 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..92
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597699
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1317:

Met Ala Arg Thr Lys Gln Thr Ala Arg Lys Ser Thr Gly Gly Lys Ala

- (2) INFORMATION FOR SEQ ID NO:1318:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 493 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..493
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597726
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1318: gaaggcatct ccaatctect egeceeccae ggaeetgaae egagaaeegt taeegeegee 60 gcctctcgcc tagccgccgg agaagcgccg ctcgctcagg aggacagcca tggcgccgga 120 gccggagatg atattatgaa cgagaagaat ccccgcccgc tcgacgagga tracatcgcg 180 ctcctcaaaa cctatggact ggggccgtat tcgaccagca ttaagaaggt cgagaaggag 240 atcaaggaaa tggccaagaa aatcaacgac ctctgtggga taaaggagtc tgatacaggg 300 ctggctccac ctagccagtg ggatttggtt tcagataagc agatgatgca agaagaacaa 360 ccgttacaag tagcaagatg tacaaagatt ataagcccta atacggatga tgccaaatat 420 480 gttataaatg traaacaaat tgcaaagttt gtggtcggat tgggagataa ggtctctcca actgatatcg agg
- (2) INFORMATION FOR SEQ ID NO:1319:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 164 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..164
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597727
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1319: Glu Gly Ile Ser Asn Leu Leu Ala Pro His Gly Pro Glu Pro Arg Thr 10 5 Val Thr Ala Ala Ala Ser Arg Leu Ala Ala Gly Glu Ala Pro Leu Ala 20 25 Gln Glu Asp Ser His Gly Ala Gly Ala Gly Asp Asp Ile Met Asn Glu 45 40 Lys Asn Pro Arg Pro Leu Asp Glu Asp Xaa Ile Ala Leu Leu Lys Thr 55 60 Tyr Gly Leu Gly Pro Tyr Ser Thr Ser Ile Lys Lys Val Glu Lys Glu 75 70 Ile Lys Glu Met Ala Lys Lys Ile Asn Asp Leu Cys Gly Ile Lys Glu 85 90 Ser Asp Thr Gly Leu Ala Pro Pro Ser Gln Trp Asp Leu Val Ser Asp 105 110 Lvs Gln Met Met Gln Glu Gln Pro Leu Gln Val Ala Arg Cys Thr 120 Lys Ile Ile Ser Pro Asn Thr Asp Asp Ala Lys Tyr Val Ile Asn Xaa

- (2) INFORMATION FOR SEQ ID NO:1320:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 119 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..119
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597728
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1320:

Met Asn Glu Lys Asn Pro Arg Pro Leu Asp Glu Asp Xaa Ile Ala Leu 1 5 10 15

Leu Lys Thr Tyr Gly Leu Gly Pro Tyr Ser Thr Ser Ile Lys Lys Val 20 25 30

Glu Lys Glu Ile Lys Glu Met Ala Lys Lys Ile Asn Asp Leu Cys Gly 35 40 45

Ile Lys Glu Ser Asp Thr Gly Leu Ala Pro Pro Ser Gln Trp Asp Leu 50 60

Val Ser Asp Lys Gln Met Met Gln Glu Glu Gln Pro Leu Gln Val Ala
65 70 75 80

Arg Cys Thr Lys Ile Ile Ser Pro Asn Thr Asp Asp Ala Lys Tyr Val 85 90 95

Ile Asn Xaa Lys Gln Ile Ala Lys Phe Val Val Gly Leu Gly Asp Lys
100 105 110

Val Ser Pro Thr Asp Ile Glu

115

- (2) INFORMATION FOR SEQ ID NO:1321:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 425 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..425
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597732
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1321:

caatgagttg aggagagtgg aacatgttga ccattctaga aagtytgcag agcargctgt 60 aaaggcgatc aagggcaaag agtccggtga accagttccg gagtatgact acctgccata 120 cttctactcc cgatcattcg acctggcgtg gcaattctac ggcgacaacg tgggcgaaac 180 tatcctgttt ggcgacagtg acccacctc cagcaagcc aagttcggct cgtactggat 240 caaggacggc aaggtcttgg gcgccttcct ggagggcggg tcaccggacg agaacaaggc 300 cattgccaag gtggcgaaaa cccagccgcc ggtcgctaac ctcgaggagc tcaagaaggm 360 tggcctccag ttcgccagca agatctgaga cgstcccgtt ccatgtaaat ttccgtgtca 420

- (2) INFORMATION FOR SEQ ID NO:1322:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 128 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:

- (A) NAME/KEY: peptide
- (B) LOCATION: 1..128
- (D) OTHER INFORMATION: / Ceres Seq. ID 1597733
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1322:
- Asn Glu Leu Arg Arg Val Glu His Val Asp His Ser Arg Lys Xaa Ala 1 5 10 15
- Glu Xaa Ala Val Lys Ala Ile Lys Gly Lys Glu Ser Gly Glu Pro Val 20 25 30
- Pro Glu Tyr Asp Tyr Leu Pro Tyr Phe Tyr Ser Arg Ser Phe Asp Leu 35 40 45
- Ala Trp Gln Phe Tyr Gly Asp Asn Val Gly Glu Thr Ile Leu Phe Gly 50 55 60
- Asp Ser Asp Pro Thr Ser Ser Lys Pro Lys Phe Gly Ser Tyr Trp Ile 65 70 75 80
- Lys Asp Gly Lys Val Leu Gly Ala Phe Leu Glu Gly Gly Ser Pro Asp
- Glu Asn Lys Ala Ile Ala Lys Val Ala Lys Thr Gln Pro Pro Val Ala 100 105 110
- Asn Leu Glu Glu Leu Lys Lys Xaa Gly Leu Gln Phe Ala Ser Lys Ile 115 120 125
- (2) INFORMATION FOR SEQ ID NO:1323:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 107 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..107
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597734
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1323:
- Met Thr Thr Cys His Thr Ser Thr Pro Asp His Ser Thr Trp Arg Gly $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$
- Asn Ser Thr Ala Thr Trp Ala Lys Leu Ser Cys Leu Ala Thr Val $20 \hspace{1cm} 25 \hspace{1cm} 30$
- Thr Pro Pro Pro Ala Ser Pro Ser Ser Ala Arg Thr Gly Ser Arg Thr 35 40 45
- Ala Arg Ser Trp Ala Pro Ser Trp Arg Ala Gly His Arg Thr Arg Thr 50 55 60
- Arg Pro Leu Pro Arg Trp Arg Lys Pro Ser Arg Arg Ser Leu Thr Ser 65 70 75 80
- Xaa Pro Phe His Val Asn Phe Arg Val Ile Val 100 105
- (2) INFORMATION FOR SEQ ID NO:1324:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 430 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..430
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597741
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1324:
- ataaaaccct actaaccgcg tctcctctcc agcgcccgcc gtcgccgccg ccctcctctt

ggtcccgccg tccgtcgagg tcatcatggt gagggtcagt gtgctcaacg atgcgctcaa 120 gtccatgtac aatgcagaga agaggggcaa gaggcaggtc atgatcaggc cgtcgtccaa 180 ggtgatcatc aagttcctga cggtcatgca gcgtcatgga tacattggcg agttcgagta 240 cgtggatgac cacagagctg ggaagattgt ggtggaactg aacggcagac taaacaaatg 300 cggtgtaatt agccccggt ttgatgttgg ggtaaaggaa atcgaaggct ggactgcgag 360 gctgcttccg tctcgccagt ttggctatat cgtgcttaca acctctgcgg gaattatgga 420 tcacgaggag

- (2) INFORMATION FOR SEQ ID NO:1325:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 115 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..115
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597742
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1325:

Met Val Arg Val Ser Val Leu Asn Asp Ala Leu Lys Ser Met Tyr Asn $1 \hspace{1.5cm} 5 \hspace{1.5cm} 10 \hspace{1.5cm} 15$

Ala Glu Lys Arg Gly Lys Arg Gln Val Met Ile Arg Pro Ser Lys 20 25 30

Val Ile Ile Lys Phe Leu Thr Val Met Gln Arg His Gly Tyr Ile Gly 35 40 45

Glu Phe Glu Tyr Val Asp Asp His Arg Ala Gly Lys Ile Val Val Glu 50 55 60

Leu Asn Gly Arg Leu Asn Lys Cys Gly Val Ile Ser Pro Arg Phe Asp 65 70 75 80

Val Gly Val Lys Glu Ile Glu Gly Trp Thr Ala Arg Leu Leu Pro Ser 85 90 95

Arg Gln Phe Gly Tyr Ile Val Leu Thr Thr Ser Ala Gly Ile Met Asp 100 105 110

His Glu Glu

115

- (2) INFORMATION FOR SEQ ID NO:1326:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 102 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..102
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597743
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1326:

Met Tyr Asn Ala Glu Lys Arg Gly Lys Arg Gln Val Met Ile Arg Pro
1 10 15

Ser Ser Lys Val Ile Ile Lys Phe Leu Thr Val Met Gln Arg His Gly

Tyr Ile Gly Glu Phe Glu Tyr Val Asp Asp His Arg Ala Gly Lys Ile
35 40 45

Val Val Glu Leu Asn Gly Arg Leu Asn Lys Cys Gly Val Ile Ser Pro
50 55 60

Arg Phe Asp Val Gly Val Lys Glu Ile Glu Gly Trp Thr Ala Arg Leu 65 70 75 80

Leu Pro Ser Arg Gln Phe Gly Tyr Ile Val Leu Thr Thr Ser Ala Gly
85 90 95

Ile Met Asp His Glu Glu

- Attorney Docket No. 750-1237P Client Docket No. 46.003 Table 2 Page 792 (2) INFORMATION FOR SEQ ID NO:1327: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 90 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..90 (D) OTHER INFORMATION: / Ceres Seq. ID 1597744 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1327: Met Ile Arg Pro Ser Ser Lys Val Ile Ile Lys Phe Leu Thr Val Met 1.0 Gln Arg His Gly Tyr Ile Gly Glu Phe Glu Tyr Val Asp Asp His Arg 25 20 Ala Gly Lys Ile Val Val Glu Leu Asn Gly Arg Leu Asn Lys Cys Gly 40 Val Ile Ser Pro Arg Phe Asp Val Gly Val Lys Glu Ile Glu Gly Trp 55 Thr Ala Arg Leu Leu Pro Ser Arg Gln Phe Gly Tyr Ile Val Leu Thr 70 75 Thr Ser Ala Gly Ile Met Asp His Glu Glu 85 (2) INFORMATION FOR SEQ ID NO:1328: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 378 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: -(B) LOCATION: 1..378 (D) OTHER INFORMATION: / Ceres Seq. ID 1597745 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1328: atcccacttc cacccaaatc caccaccgcc ataccagcgt gtccgcgatt ccgtcqagtt 60 agaccagtcg agagagcagt cccagacgag aagaggtgaa tatgagttcc ggcggcggca 120 ggggcaagcc caaggggtcc aaggctgtgt cgcggtcgac caaggctggg ctgcagttcc 180 ccgtcggccg catcgcgcgc tacctcaagg cgggcaagta cgccgagcgc gtcggcggtg 240 gtgcgcccgt ctacctctcc gccgtcctcg agtatctcgc ggcagaggtg ttggaactgg 300 cgggcaacgc ggcgcgcac aacaagaaga accgcatcgt gccgcgccac atccagctcg 360 ccqtqcqcaa cqacqagg (2) INFORMATION FOR SEQ ID NO:1329: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 125 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..125
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597746 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1329:

 Ser His Phe His Pro Asn Pro Pro Pro Pro Tyr Gln Arg Val Arg Asp

 1
 5
 5
 10
 15

 Ser Val Glu Leu Asp Gln Ser Arg Glu Gln Ser Gln Thr Arg Arg Gly
 25
 30

Glu Tyr Glu Phe Arg Arg Gln Gly Gln Ala Gln Gly Val Gln Gly
35 40 45

Cys Val Ala Val Asp Gln Gly Trp Ala Ala Val Pro Arg Arg Pro His 50 60

Arg Ala Leu Pro Gln Gly Gly Gln Val Arg Arg Ala Arg Arg Arg Trp 65 70 75 80

Cys Ala Arg Leu Pro Leu Arg Arg Pro Arg Val Ser Arg Gly Arg Gly
85 90 95

Val Gly Thr Gly Gly Gln Arg Gly Ala Arg Gln Gln Glu Glu Pro His 100 105 110

Arg Ala Ala Pro His Pro Ala Arg Arg Ala Gln Arg Arg 115 120 125

- (2) INFORMATION FOR SEQ ID NO:1330:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 92 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..92
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597747
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1330:

Met Ser Ser Gly Gly Gly Arg Gly Lys Pro Lys Gly Ser Lys Ala Val 1 5 10 15

Ser Arg Ser Thr Lys Ala Gly Leu Gln Phe Pro Val Gly Arg Ile Ala 20 25 30

Arg Tyr Leu Lys Ala Gly Lys Tyr Ala Glu Arg Val Gly Gly Gly Ala 35 40 45

Pro Val Tyr Leu Ser Ala Val Leu Glu Tyr Leu Ala Ala Glu Val Leu 50 55 60

Glu Leu Ala Gly Asn Ala Ala Arg Asp Asn Lys Lys Asn Arg Ile Val 65 70 75 80

Pro Arg His Ile Gln Leu Ala Val Arg Asn Asp Glu 85 90

- (2) INFORMATION FOR SEQ ID NO:1331:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 456 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..456
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597748
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1331:

ctctagaacc ctagcgcbnc gcccgtcctg ccgccgcc agccgccatc gcacggccat 60 gggtatcgat ctcgtagccg gcggccgcaa caagaakacc aagcgcacgg cgcccaagtc 120 cgacgatgtc tacctcaagc tcctcgtcaa gctctaccgc ttccttgtgc gcaggacaaa 180 gagcaacttc aacgccgtga tcctcaagcg cctcttcatg agcaagacca accgcccgcc 240 gctctcggat gcgccgcctc gtcaggttca tggaggggaa gggtgatcag atcgccgtga 120 gctctcggat gcgccgccc gtcaggttca tggaggggaa gggtgatcag atcgccgtga 300 ttgtgggcac cgtgaccgac gacaagagga tcagcgaggt gccggcgatg aaggtctgcg 360 cgctcaggtt cactgagaca gcgagggcca ggatcatcaa tgctggcgga gagtgcctca 420 ccttcqacca gctcgcgct cgtgccccac ttgggc

- (2) INFORMATION FOR SEQ ID NO:1332:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 94 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide

- (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..94
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597749
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1332:
- Ser Arg Thr Leu Ala Xaa Arg Pro Ser Cys Arg Arg Arg Ser Arg His
 1 10 15
- Arg Thr Ala Met Gly Ile Asp Leu Val Ala Gly Gly Arg Asn Lys Xaa 20 25 30
- Thr Lys Arg Thr Ala Pro Lys Ser Asp Asp Val Tyr Leu Lys Leu Leu 35 40 45
- Val Lys Leu Tyr Arg Phe Leu Val Arg Arg Thr Lys Ser Asn Phe Asn 50 55 60
- Ala Val Ile Leu Lys Arg Leu Phe Met Ser Lys Thr Asn Arg Pro Pro 65 70 75 80
- Leu Ser Asp Ala Pro Pro Arg Gln Val His Gly Gly Glu Gly 85 90
- (2) INFORMATION FOR SEQ ID NO:1333:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 75 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..75
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597750
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1333:
- Met Gly Ile Asp Leu Val Ala Gly Gly Arg Asn Lys Xaa Thr Lys Arg 1 5 10 15
- Thr Ala Pro Lys Ser Asp Asp Val Tyr Leu Lys Leu Leu Val Lys Leu 20 25 30
- Tyr Arg Phe Leu Val Arg Arg Thr Lys Ser Asn Phe Asn Ala Val Ile $35 \hspace{1cm} 40 \hspace{1cm} 45$
- Leu Lys Arg Leu Phe Met Ser Lys Thr Asn Arg Pro Pro Leu Ser Asp 50 55 60
- Ala Pro Pro Arg Gln Val His Gly Gly Glu Gly 65 70 75
- (2) INFORMATION FOR SEQ ID NO:1334:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 69 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..69
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597751
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1334:
- Met Arg Arg Leu Val Arg Phe Met Glu Gly Lys Gly Asp Gln Ile Ala 1 5 10 15
- Val Ile Val Gly Thr Val Thr Asp Asp Lys Arg Ile Ser Glu Val Pro 20 25 30
- Ala Met Lys Val Cys Ala Leu Arg Phe Thr Glu Thr Ala Arg Ala Arg 35 40 45
- Ile Ile Asn Ala Gly Gly Glu Cys Leu Thr Phe Asp Gln Leu Ala Leu 50 55 60
- Arg Ala Pro Leu Gly

60

120

180

240

(2) INFORMATION FOR SEQ ID NO:1335: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 294 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: DNA (genomic) (ix) FEATURE: (A) NAME/KEY: -(B) LOCATION: 1..294 (D) OTHER INFORMATION: / Ceres Seq. ID 1597752 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1335: gttgttttgt tcttgcggaa acgagacgag actaggcgac gcgasgtyag ggagtcgagg aggategggg eteaggeaag atgaacagga agecaggaga etgggaetge agggegtgee ancacctcaa cttcagccgc cgagacatat gccagckctg tagcgagcca cgtggagttg ctgatcgtgg cagtggcggc ggcggaggag gaggaggagg cgactacgca agctttcggt ggccgcggtg gctcctcctt cggcggcggc tttgggcgct gctggctctg acgt (2) INFORMATION FOR SEQ ID NO:1336: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 53 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..53 (D) OTHER INFORMATION: / Ceres Seq. ID 1597753 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1336: Leu Phe Cys Ser Cys Gly Asn Glu Thr Arg Leu Gly Asp Ala Xaa Xaa 10 Gly Ser Arg Gly Gly Ser Gly Leu Arg Gln Asp Glu Gln Glu Ala Arg 25 Arg Leu Gly Leu Gln Gly Val Pro Xaa Pro Gln Leu Gln Pro Pro Arg His Met Pro Xaa Leu 50 (2) INFORMATION FOR SEQ ID NO:1337: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 71 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (ix) FEATURE: (A) NAME/KEY: peptide (B) LOCATION: 1..71 (D) OTHER INFORMATION: / Ceres Seq. ID 1597754 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1337: Met Asn Arg Lys Pro Gly Asp Trp Asp Cys Arg Ala Cys Xaa His Leu Asn Phe Ser Arg Arg Asp Ile Cys Gln Xaa Cys Ser Glu Pro Arg Gly Val Ala Asp Arg Gly Ser Gly Gly Gly Gly Gly Gly Gly Gly Asp 40 Tyr Ala Ser Phe Arg Trp Pro Arg Trp Leu Leu Arg Arg Arg Leu

70 (2) INFORMATION FOR SEQ ID NO:1338:

Trp Ala Leu Leu Ala Leu Thr

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 442 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA (genomic)
- (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..442
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597755
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1338:

cttttaacct cgcgccgca tccgacgcc atctgctca ccctccacc agccgccgcc 60 gccgccgcct aaggagaaga gggtccggcc gtctccttgc cccgatggcg tccgagaaga 120 agcagtccaa cccgatgcgg gagatcaagg tgcagaagct ggtcctcaac atatccgtag 180 ggagaggcgc gaccgcctca cccgcgccgc caaggtgctg gagcagctca gcgccagacc 240 cccgtcttct ccaaggcgag gtacacggtg cgctcgttcg gcatccggcg taacgagaag 300 atcgcctgct acgtcacggt caggggcgac aaggccatgc agctgcttga gagcggcctc 360 aaggtcaagg agtacgagct gctcaggagg aacttcagcg acaccggctg cttcgggttc 420 ggcatccagg agcacatcga cc

- (2) INFORMATION FOR SEQ ID NO:1339:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 96 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..96
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597756
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1339:

Phe Asn Leu Ala Pro Pro Ser Asp Ala His Leu Leu His Pro Pro Pro $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Ser Arg Arg Arg Arg Leu Arg Arg Gly Ser Gly Arg Leu Leu 20 25 30

Ala Pro Met Ala Ser Glu Lys Lys Gln Ser Asn Pro Met Arg Glu Ile $35 \hspace{1cm} 40 \hspace{1cm} 45$

Lys Val Gln Lys Leu Val Leu Asn Ile Ser Val Gly Arg Ala Ala Thr 50 55 60

Ala Ser Pro Ala Pro Pro Arg Cys Trp Ser Ser Ser Ala Pro Asp Pro 65 70 75 80

Arg Leu Leu Gln Gly Glu Val His Gly Ala Leu Val Arg His Pro Ala 85 90 95

- (2) INFORMATION FOR SEQ ID NO:1340:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 62 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..62
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597757
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1340:

Met Ala Ser Glu Lys Lys Gln Ser Asn Pro Met Arg Glu Ile Lys Val 1 5 10 15

Gln Lys Leu Val Leu Asn Ile Ser Val Gly Arg Ala Ala Thr Ala Ser 20 25 30

Pro Ala Pro Pro Arg Cys Trp Ser Ser Ser Ala Pro Asp Pro Arg Leu

35 40 45 Leu Gln Gly Glu Val His Gly Ala Leu Val Arg His Pro Ala 50 55 60

- (2) INFORMATION FOR SEQ ID NO:1341:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 52 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..52
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597758
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1341:

Arg Ala Ala Thr Ala Ser Pro Ala Pro Pro Arg Cys Trp Ser Ser Ser 20 25 30

Ala Pro Asp Pro Arg Leu Leu Gln Gly Glu Val His Gly Ala Leu Val 35 40 45

Arg His Pro Ala 50

- (2) INFORMATION FOR SEQ ID NO:1342:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 422 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..422
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597797
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1342:

ategecege gaacectage teegeaact geegeege eteceaeget egeageeatg 60 cegeceaage tegaceegte geaggtggte gaggtgtteg teegtgtgae eggeggegg 120 gteggegeae gtettegetg geececaaga teggeeeget eggtettee eceaagaaga 180 teggagagga categeeaag gagaeggea aggaetggaa gggeeteege gteaeegtea 240 ageteaeegt eeagaacegg eageeaaggt eteegtegte eeeteegeeg eggegetegt 240 cateaaggeg eteaaggag eegeaagga eegeaagga gteaagaaca teaageacag egneaacate ageetegae aegteatega gategeeagg aecatgagge aecatgagge aecatgagge gg

- (2) INFORMATION FOR SEQ ID NO:1343:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 140 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..140
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597798
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1343:
- Ile Ala Pro Arg Thr Leu Ala Pro Ala Thr Ala Ala Ala Thr Pro Thr

 10 15
- Leu Ala Ala Met Pro Pro Lys Leu Asp Pro Ser Gln Val Val Glu Val
 20 25 30
- Phe Val Arg Val Thr Gly Gly Glu Val Gly Ala Arg Leu Arg Trp Pro
 35

Pro Arg Ser Ala Arg Ser Val Phe Pro Pro Arg Arg Ser Glu Arg Thr

Ser Pro Arg Arg Arg Pro Arg Thr Gly Arg Ala Ser Ala Ser Pro Ser 65 70 75 80

Ser Ser Pro Ser Arg Thr Gly Ser Gln Gly Leu Arg Arg Pro Leu Arg 85 90 95

Arg Gly Ala Arg His Gln Gly Ala Gln Gly Ala Arg Glu Gly Pro Gln 100 105 110

Glu Gly Gln Glu His Gln Ala Gln Xaa Gln His Gln Pro Arg Arg 115 120 125

His Arg Asp Arg Gln Asp His Glu Ala Gln Val His 130 $\,$ 135 $\,$ 140

- (2) INFORMATION FOR SEQ ID NO:1344:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 134 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..134
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597799
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1344:

Arg Pro Ala Asn Pro Ser Ser Arg Asn Cys Arg Arg His Ser His Ala 1 10 15

Arg Ser His Ala Ala Gln Ala Arg Pro Val Ala Gly Gly Gly Val
20 25 30

Arg Pro Cys Asp Arg Arg Gly Arg Arg Thr Ser Ser Leu Ala Pro
35 40 45

Lys Ile Gly Pro Leu Gly Leu Ser Pro Lys Lys Ile Gly Glu Asp Ile

Ala Lys Glu Thr Ala Lys Asp Trp Lys Gly Leu Arg Val Thr Val Lys 65 70 75 80

Leu Thr Val Gln Asn Arg Gln Pro Arg Ser Pro Ser Ser Pro Pro Pro 85 90 95

Arg Arg Ser Ser Ser Arg Arg Ser Arg Ser Gln Arg Gly Thr Ala Arg
100
105
110

Arg Ser Arg Thr Ser Ser Thr Ala Xaa Thr Ser Ala Ser Thr Thr Ser 115 120 125

Ser Arg Ser Pro Gly Pro 130

- (2) INFORMATION FOR SEQ ID NO:1345:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 121 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..121
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597800

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1345:

Met Pro Pro Lys Leu Asp Pro Ser Gln Val Val Glu Val Phe Val Arg
1 5 10 15

Val Thr Gly Gly Glu Val Gly Ala Arg Leu Arg Trp Pro Pro Arg Ser 20 25 30

Ala Arg Ser Val Phe Pro Pro Arg Arg Ser Glu Arg Thr Ser Pro Arg 35 40 45

Arg Arg Pro Arg Thr Gly Arg Ala Ser Ala Ser Pro Ser Ser Pro

60

120

180

240

300

360

420

480

- (2) INFORMATION FOR SEQ ID NO:1346:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 483 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (ix) FEATURE:
 - (A) NAME/KEY: -
 - (B) LOCATION: 1..483
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597805
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1346: aaaaccctag ccgcctgcyt ctcctcctcc agagctcgtc gccaccggcc accgccaccc tcgatcacgc cgtccgtcga ggtaaggtca gcatggtgag ggtcagtgtg ctcaacgatg cgctcaagtc catgtacaac gctgagaaga tcggcaagag gcaggtcatg atcaggccgt cgtccaaggt catcatcaag ttcctgacgg tcatgcagcg ccacggctac attggagagt tcgagtacgt tgatgaccac cgatcggca agatcgtggt cgaactcaac gggaggctga

acaagtgegg egteateage eetegetttg atateggegt gaaagacatt gagggatgga etgeaagetg etecegteea ggeagttegg atacategte eteacaactt eggeaggeat

catggaccac gaggaggece geeggaagag egtaggagge aaggttetag gtttetteta

- (2) INFORMATION FOR SEQ ID NO:1347:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 150 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..150
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597806
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1347:
- Asn Pro Ser Arg Leu Xaa Leu Leu Leu Gln Ser Ser Pro Pro Ala 1 5 10 15
- Thr Ala Thr Leu Asp His Ala Val Arg Arg Gly Lys Val Ser Met Val
 20 25 30
- Arg Val Ser Val Leu Asn Asp Ala Leu Lys Ser Met Tyr Asn Ala Glu 35 40 45
- Lys Ile Gly Lys Arg Gln Val Met Ile Arg Pro Ser Ser Lys Val Ile
 50 55 60
- Ile Lys Phe Leu Thr Val Met Gln Arg His Gly Tyr Ile Gly Glu Phe 65 70 75 80
- Glu Tyr Val Asp Asp His Arg Ser Gly Lys Ile Val Val Glu Leu Asn 85 90 95
- Gly Arg Leu Asn Lys Cys Gly Val Ile Ser Pro Arg Phe Asp Ile Gly 100 105 110
- Val Lys Asp Ile Glu Gly Trp Thr Ala Ser Cys Ser Arg Pro Gly Ser 115 120 125
- Ser Asp Thr Ser Ser Ser Gln Leu Arg Gln Ala Ser Trp Thr Thr Arg 130 135 140
- Arg Pro Ala Gly Arg Ala

Attorney Docket No. 50-1237P Client Docket No. 6.003

145 150

- (2) INFORMATION FOR SEQ ID NO:1348:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 120 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..120
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597807
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1348:

Met Val Arg Val Ser Val Leu Asn Asp Ala Leu Lys Ser Met Tyr Asn

5 10 15

Ala Glu Lys Ile Gly Lys Arg Gln Val Met Ile Arg Pro Ser Ser Lys 20 25 30

Val Ile Ile Lys Phe Leu Thr Val Met Gln Arg His Gly Tyr Ile Gly 35 40 45

Glu Phe Glu Tyr Val Asp Asp His Arg Ser Gly Lys Ile Val Val Glu 50 55 60

Leu Asn Gly Arg Leu Asn Lys Cys Gly Val Ile Ser Pro Arg Phe Asp 65 70 75 80

Ile Gly Val Lys Asp Ile Glu Gly Trp Thr Ala Ser Cys Ser Arg Pro
85 90 95

Gly Ser Ser Asp Thr Ser Ser Ser Gln Leu Arg Gln Ala Ser Trp Thr 100 105 110

Thr Arg Arg Pro Ala Gly Arg Ala 115 120

- (2) INFORMATION FOR SEQ ID NO:1349:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 107 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: peptide
 - (B) LOCATION: 1..107
 - (D) OTHER INFORMATION: / Ceres Seq. ID 1597808
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1349:

Ser Ser Lys Val Ile Ile Lys Phe Leu Thr Val Met Gln Arg His Gly 20 25 30

Tyr Ile Gly Glu Phe Glu Tyr Val Asp Asp His Arg Ser Gly Lys Ile 35 40 45

Val Val Glu Leu Asn Gly Arg Leu Asn Lys Cys Gly Val Ile Ser Pro 50 60

Arg Phe Asp Ile Gly Val Lys Asp Ile Glu Gly Trp Thr Ala Ser Cys 65 70 75 80

Ser Arg Pro Gly Ser Ser Asp Thr Ser Ser Ser Gln Leu Arg Gln Ala 85 90 95

Ser Trp Thr Thr Arg Arg Pro Ala Gly Arg Ala 100 105

- (2) INFORMATION FOR SEQ ID NO:1350:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 133 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear